

PERSPECTIVE



Polo-like kinase 3, hypoxic responses, and tumorigenesis

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ABSTRACT

The cellular hypoxic response contributes to cell transformation and tumor progression. Hypoxia-inducible factor 1 (HIF-1) is a key transcription factor that mediates transcription of genes whose products are essential for cellular adaptation to hypoxia. The activity of HIF-1 is largely regulated by the abundance of its alpha subunit (HIF-1 α), which is primarily regulated by an oxygen-dependent and ubiquitin/proteasome-mediated degradation process. The HIF-1 α protein level is also regulated by protein kinases through phosphorylation. Polo-like kinase 3 (Plk3) is a serine/threonine protein kinase with a tumor suppressive function. Plk3 phosphorylates and destabilizes HIF-1 α . Plk3 also phosphorylates and stabilizes PTEN, a known regulator of HIF-1 α stability via the PI3K pathway. Our latest study showed that the Plk3 protein is suppressed by hypoxia or nickel treatment via the ubiquitin/proteasome system. We discovered that Seven in Absentia Homologue 2 (SIAH2) is the E3 ubiquitin ligase of Plk3 and that Plk3 in turn destabilizes SIAH2. Given the role of SIAH2 in promoting stability of HIF-1 α , our work reveals a novel mutual regulatory mechanism between Plk3 and SIAH2, which may function to fine-tune the cellular hypoxic response. Here we discuss the role of Plk3 in the hypoxic response and tumorigenesis in light of these latest findings.

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The hypoxic response

The cellular response to hypoxia (the hypoxic response) is a complex intracellular signaling network that coordinates the biological activities in response to low oxygen tension.^{1,2} The central players of this response are hypoxia inducible factors (HIFs),^{1,2} with HIF-1 being the most important and best characterized.^{1,3} As a transcription factor, HIF-1 mediates expression of a group of HIF-response genes, such as the vascular endothelial growth factor (VEGF) and the glucose transporter GLUT1, by binding to the HIF response element (HRE) on the promoters of these genes.¹⁻³ Expression of these genes triggers further signaling cascades leading to profound biological changes in the cell. These changes include metabolic alterations that stimulate angiogenesis and promote cell survival, which are required for adaptation to hypoxia.¹⁻³

The overall cellular activity of HIF-1 is primarily dictated by the abundance of its α subunit (HIF-1 α), which is regulated mainly at the post-translational level.¹⁻³ HIF-1 α is inducible under hypoxia whereas the beta subunit (HIF-1 β or ARNT) is constitutively expressed.¹⁻³ Under normoxia, HIF-1 α is degraded through oxygen-dependent hydroxylation and the ubiquitin proteasome system mediated by prolyl hydroxylases (PHDs) and Von Hippel-Lindau Factor (pVHL), respectively.¹⁻³ Low oxygen tension reduces hydroxylation and slows degradation of HIF-1 α , which lead to higher overall HIF-1 activity in the cell.¹⁻³ The stability of HIF-1 α is also regulated by protein kinases. Phosphorylation by a number of protein kinases,

including ERKs, GSK3 β , and Plk3, has been shown to alter HIF-1 α stability and/or localization.⁴⁻⁹

The hypoxic response contributes to both tumorigenesis and tumor progression.^{3,10,11} An overactive hypoxic response pathway, often manifested as elevated cellular levels of HIF-1 α , is a common feature of human cancers.^{3,11} This characteristic is essential for the survival of rapidly proliferating tumor cells that frequently face oxygen and/or nutrient restrictions due to increase in tumor masses. Elevated hypoxic responses promote survival of tumor cells by triggering angiogenesis which supplies nutrients and oxygen as well as by reprogramming the cellular metabolism, all of which are essential for cellular adaptation to the hypoxic condition.^{1-3,11} Given the importance of the hypoxic response in cancer, inhibition of the hypoxic pathway and/or tumor angiogenesis is considered an important strategy of cancer therapy.^{12,13}

Polo-like kinase 3

Polo-like kinase 3 (Plk3) is one of the 5 mammalian members (Plk1-5) of an evolutionarily conserved family of serine/threonine protein kinases, which share significant amino acid sequence homology.¹⁴⁻¹⁷ All Plks have a highly conserved kinase domain (KD) at the amino-terminus and a polo box domain (PBD) at the carboxyl-terminus.¹⁴⁻¹⁷ The KD of Plks confers catalytic activity whereas the PBD is important for their subcellular localization and the substrate recognition.¹⁵⁻¹⁷ Plk1 is the best studied member of the Plk family with a well-defined

and critical role in cell cycle progression.¹⁵⁻¹⁷ Plk4 has an important role in centrosome dynamic during the cell cycle.¹⁴⁻¹⁷ Plk5 has a truncated KD rendering it kinase-deficient.¹⁴⁻¹⁹ Plk5 functions to suppress cell cycle progression, mediate neuron differentiation, and suppress glioblastoma.^{18,19} The expression of Plk5 appears to be restricted to the brain.¹⁵⁻¹⁹ The functions of Plk2 and Plk3 seem to be more diverse and not restricted to cell cycle progression.^{14-17,19} Importantly, all members of the Plk kinase family have close association with tumorigenesis and tumor progression.^{14-17,19-22}

PLK3 is considered an immediate early response gene whose mRNA level is inducible by mitogenic stimulation.^{15,21} Interestingly, the level of Plk3 protein is quite constant in mitogen-stimulated cells throughout the cell cycle,^{15,21} although it has also been reported that the Plk3 protein level does oscillate during the cell cycle.^{23,24} The kinase activity of Plk3, on the other hand, appears to oscillate during the cell cycle and is regulated by a variety of stress conditions, including genotoxic insults, hypoxia, and osmotic stresses.²⁵⁻²⁸ The functional profile of Plk3 is apparently rather diverse. Earlier work indicates that Plk3 is involved in multiple phases of cell cycle progression, including G1/S transition, mitosis, DNA replication, Golgi fragmentation, and centrosomal functions.^{15,21} Later studies revealed additional functions of Plk3 in stress responses.^{8,25,27,28} Despite the functional significance of Plk3 discovered in various cell-based studies, *PLK3* null mice are rather normal and fertile.^{9,29} However, these mice tend to be slightly larger and more prone to spontaneous tumor development later in the life.⁹ This is in sharp contrast with the embryonic lethal phenotypes as result of deletion of Plk1 or Plk4.^{30,31} The lack of a significant adverse phenotype suggests that cellular functions of Plk3, particularly those associated with cell cycle regulation, can be compensated by other members of the Plk family and therefore largely dispensable. Functional complementation studies by us and others show that both Plk3 and Plk1 are capable of rescuing CDC5 (Plk of budding yeast)-deficiency in budding yeast.^{32,33} It appears that divergent evolution eventually leads to new functions of Plk3 in higher animals despite its conserved functions in yeast. Thus, regulation of stress responses rather than normal cell cycle progression could be the primary function of Plk3 in mammals.

Plk3 expression is reduced in many human malignancies, including those in the lung, head and neck, colon, kidney, liver, stomach, and rectum.^{15,21} Expression of Plk3 mRNA and protein is also significantly deregulated in human melanoma cell lines and tissues.¹⁵ Plk3 mRNA was found to be significantly downregulated in a majority of more than a dozen human lung carcinoma samples, apparently as a result of reduced *PLK3* transcription.³⁴ These data suggest that reduced Plk3 expression may be associated with tumor development. This notion is supported by the observation that although polymorphisms were identified in 40 lung tumor cell lines, no missense or nonsense mutations were found in Plk3.³⁵ These previous observations and the finding that *PLK3* null mice are prone to developing tumors in several organs later in the life indicate a tumor suppressive role of Plk3 and that reduced expression is likely the main mechanism that associates Plk3 with increased tumorigenesis.

Regulation of HIF-1 α by Plk3 through direct phosphorylation

The implication of Plk3 in the cellular hypoxic response was initially revealed in a genetic study showing that *PLK3* null mice exhibited an increased tumor incidence later in the life and that the tumors developed in these mice were often larger and more vasculated than those from the wild type animals.⁹ Biochemical analysis showed that murine embryonic fibroblasts (MEFs) from *PLK3* null mice express a much elevated level of HIF-1 α in response to hypoxia or nickel, a hypoxia mimic.^{8,9} Furthermore, ectopically expressed Plk3 suppresses nuclear accumulation of HIF-1 α in HeLa cells.⁹ Inhibition of HIF-1 α nuclear translocation appears to be dependent on the kinase activity of Plk3 as overexpression of the Plk3 kinase domain was sufficient to suppress HIF-1 α accumulation in the nucleus under hypoxic conditions.⁹ Consistently, expression of VEGF-A, a major HIF-1 α response protein, was also higher in *PLK3* null MEFs.⁹ These results suggest a possible direct regulation of HIF-1 α by Plk3. Follow up studies using *in vitro* kinase assay in combination with mass spectrometry confirmed that Plk3 phosphorylates HIF-1 α at two evolutionarily conserved serine residuals: Ser-576 and Ser-657.⁸ Ser-576 is located within the oxygen-dependent degradation domain (ODDD) whereas Ser-657 residues immediate downstream of the nuclear export signal (NES) of HIF-1 α ,⁸ suggesting that Plk3 may regulate degradation and nuclear export of HIF-1 α . Further experimentation confirmed that phosphorylation of these residuals reduces the stability of HIF-1 α in a hydroxylation- and pVHL-independent manner.⁸

Previous work demonstrated that ERK MAP kinases phosphorylate HIF-1 α at residues Ser-641 and Ser-643 (both are within NES), through which promotes translocation of HIF-1 α from the cytoplasm to the nucleus.^{6,7} Glycogen synthase kinase 3 β (GSK3 β) phosphorylates HIF-1 α at three serine residues (Ser-551, Ser-555, and Ser-589) located within ODDD,⁴ through which enhances HIF-1 α degradation in a pVHL-independent manner.^{4,36} The discovery that Plk3 regulates HIF-1 α added one more kinase to the short list of protein kinases that directly regulate HIF-1 α through direct phosphorylation.

Regulation of HIF-1 α by Plk3 through PTEN

Phosphatase and tensin homologue (PTEN) is an important tumor suppressor that inhibits the phosphatidylinositol 3-kinases kinase (PI3K) signaling pathway by dephosphorylating the phosphoinositides.^{37,38} Activation of the PI3K pathway leads to an elevated AKT activity.³⁸ AKT may increase the HIF-1 α protein level by activating mTOR or inhibiting GSK3 β , which regulate the protein synthesis and stability of HIF-1 α , respectively.^{4,12,39-41}

PTEN can be phosphorylated by a number of kinases.⁴²⁻⁴⁹ Phosphorylation of PTEN can affect its activity and/or stability⁴²⁻⁵². The sites of phosphorylation on PTEN are concentrated at the C-terminal region of the protein,⁴²⁻⁴⁹ the regulatory domain of PTEN.^{42,53,54} Phosphorylation of PTEN by Plk3 was discovered based on the observation that *PLK3* null MEFs exhibited reduced levels of the PTEN protein.⁴³ *In vitro* kinase assays followed by mass spectrometry identified Thr-366 and Ser-370 at the C-

terminal region of PTEN as the phosphorylation targets of Plk3.⁴³ These two sites were further confirmed using a phospho-specific antibody that recognized p-Thr-366 and p-Ser-370.⁴³ Phosphorylation of these two residues enhances the stability of PTEN, consistent with the reduced PTEN protein level in *PLK3* null MEFs.⁴³ Thus, phosphorylation of these two sites by Plk3 may stabilize PTEN and lead to an increased overall PTEN activity in the cell. Given the known effect of the PI3K pathway on HIF-1 α stability and that PTEN is a negative regulator of the PI3K pathway, it is conceivable that Plk3 may affect HIF-1 α stability indirectly through the PI3K signaling pathway.

Regulation of HIF-1 α through mutual regulation between Plk3 and SIAH2

Our most recent work has added additional complexity to the regulation of HIF-1 α /the hypoxic response by Plk3. A recent effort to understand the effects of hypoxia and nickel on Plk3 expression reveals that the Plk3 protein is suppressed by hypoxia or nickel through the ubiquitin proteasome system.⁵⁵ Seven in Absentia Homologue 2 (SIAH2), a RING finger E3 ubiquitin ligase, was identified to catalyze ubiquitination of Plk3 and to promote Plk3 degradation.⁵⁵ SIAH2 apparently interacts with Plk3 through two domains that closely resemble the consensus SIAH2 binding motif.⁵⁵⁻⁵⁷ One of these domains is located within the KD of Plk3 whereas the other one resides slightly N-terminal of the PBD.⁵⁵ The domain near the PBD seems to be the main site for the interaction and Plk3 degradation.⁵⁵ SIAH2 has been shown to be activated and induced by hypoxia and in turn mediates the ubiquitination and degradation of PHDs.^{58,59} SIAH2 appears to regulate Plk3 in a similar fashion. Given that both PHDs and Plk3 negatively regulate the stability of HIF-1 α , SIAH2 may regulate HIF-1 α via both PHDs and Plk3. More interestingly, Plk3 also destabilizes SIAH2 in a kinase activity-dependent manner.⁵⁵ Thus, a mutual regulatory mechanism exists between Plk3 and SIAH2, which may function to fine-tune the HIF-1 signaling.

USP28, a deubiquitinase that suppresses the stability of MYC and HIF-1 α ,^{36,60,61} also appears to indirectly contribute to suppression of Plk3 by hypoxia and nickel.⁵⁵ It has been shown that USP28 can be suppressed by nickel via HIF-, the ubiquitin-proteasome system-, and DNA methylation-dependent mechanisms.⁶² USP28 prevents the suppression of Plk3 by nickel, suggests that suppression of Plk3 deubiquitination by USP28 in response to nickel could contribute to the elevated degradation of Plk3 by the ubiquitin-proteasome system. However, the effect of USP28 on Plk3 is likely indirect as a direct interaction between Plk3 and USP28 was not detected.⁵⁵ Of note, although USP28 has been reported to mediate HIF-1 α stability, a direct interaction between the two was also undetectable.³⁶

Taken together, the newest findings reveal a novel mutual regulatory mechanism between Plk3 and SIAH2 and support a complex role of Plk3 in regulating the cellular hypoxic response through HIF-1 α : under normoxia, Plk3 suppresses the hypoxic response by phosphorylating and destabilizing HIF-1 α and SIAH2; under hypoxia or the hypoxia-like condition induced by nickel, the level and/or activity of SIAH2 increases and the level of USP28 decreases, which suppress

the protein levels of Plk3 and PHDs; Reduced expression of Plk3 and PHDs in turn helps maintain HIF-1 α and SIAH2 proteins at higher levels. This mutual regulatory network highlights a potentially important role of Plk3 in a signaling network that functions to fine-tune the cellular hypoxic response.

Implication of the regulatory network of Plk3, SIAH2, and HIF-1 α in tumorigenesis

Evidence collected thus far has established a tumor suppressive role of Plk3 through a mechanism independent of its previously discovered functions on cell cycle regulation. Despite the observed functions of Plk3 in multiple biological processes associated with cell cycle progression at the cellular level, the *PLK3* null mouse is largely normal.⁹ Discernable phenotypes of *PLK3* null mice are the slightly larger sizes and the higher tendency of developing highly vasculated tumors in multiple organs.⁹ These phenotypes are consistent with the findings on the regulation of PTEN and HIF-1 pathways by Plk3 as these pathways are known to regulate cell growth, cell survival, and tumor angiogenesis.^{1-3,39,63} *In vivo* data also imply that many Plk3 functions described earlier based on molecular and cellular studies are largely dispensable for normal mouse physiology. This is likely a result of the functional redundancy of Plk3 with other members of the Plk family. However, the phenotypes of *PLK3* null mice on tumor burden and the hypoxic response strongly suggest that Plk3 may suppress spontaneous tumorigenesis and tumorigenesis induced by carcinogens, particularly those mimicking hypoxia, such as nickel compounds and other metal carcinogens.

SIAH2 is considered an oncogene in multiple tissues, including the lung.^{59,64} Elevated expression of SIAH2 has been detected in lung cancers.^{64,65} SIAH2 promotes tumorigenesis through the Ras signaling pathway by targeting the Ras inhibitor Sprouty 2 for degradation as well as through the hypoxic response pathway.^{58,64} The finding that Plk3 destabilizes SIAH2 in a kinase activity-dependent manner highlights an additional mechanism underlying the role of Plk3 in tumorigenesis. In conclusion, recent studies have provided new mechanistic insights on how Plk3 may contribute to tumorigenesis and tumor progression (Figure 1).

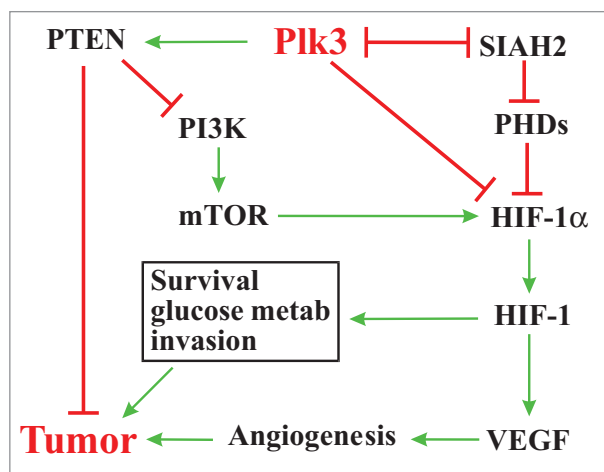


Figure 1. Plk3, hypoxic responses, and tumorigenesis.

Perspectives

Unlike Plk1, the prototype of the mammalian Plk kinase family, the biology of Plk3 and its role in tumor biology is much less studied and understood. Recent discoveries on the new functions of Plk3 in the HIF pathway have shed fresh light on the importance and mechanisms of this protein in tumorigenesis and tumor progression. While Plk1 has been viewed as an attractive target for cancer therapy based on its well defined functions in cell cycle progression, the potential of Plk3 in this regard has not been fully appreciated. Given the role of Plk3 in the hypoxic response, it is conceivable that this protein kinase can be a very significant player in tumorigenesis and thus serve as a therapeutic target and/or tumor biomarker. Further studies on the biological significance and detailed mechanisms of Plk3 in regulating the hypoxic pathway and tumorigenesis, particularly *in vivo*, are highly warranted.

Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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