

Correlation of histopathological high risk factors with Polo-Like Kinase 1 (PLK1) in retinoblastoma

Lata Singh

All India Institute of Medical Sciences, India

Abstract

Background: Retinoblastoma remains a therapeutic challenge for pediatric oncologists. Polo-like kinases (Plks), a family of conserved serine/threonine kinases, are important regulators of cell cycle progression for maintaining DNA integrity. Expression of PLK1 in retinoblastoma has not been studied so far and so its role remains unclear till now. Retrospective analyses of 30 primary enucleated retinoblastoma cases over a period of 2010-2011 were included in this study. PLK1 protein expression was performed by immunohistochemistry in formalin fixed retinoblastoma specimens. Cytoplasmic staining was graded as weak/negative (1+), moderate (2+) and strong (3+). PLK1 expression was correlated with tumour differentiation and histopathological high risk factors. The patients were followed up for one year. Out of 30 eyes, 22 were poorly differentiated retinoblastoma and 8 were well differentiated retinoblastoma. Extensive necrosis and calcification was found in 53.3% and 20% respectively. Histopathologically, 11 cases had massive choroidal invasion, 14 optic nerve invasion, 2 each scleral and anterior chamber, 3 with iris and ciliary body invasion. PLK1 expression was observed in 22/30 (73.3%) cases. Of the 22 cases, there were total of 17 cases in which more than one histopathological high risk factor was present. However, no statistically significant difference was seen between PLK1 expression and histopathological high risk factors. All the patients were alive without any recurrences until last follow up. Overexpression of PLK1 was observed in cases with one or more histopathological high risk factor in retinoblastoma. These findings suggest that PLK1 may be useful as prognostic marker in patients with Retinoblastoma.

Keywords

Histopathologically, PLK1

Background

Retinoblastoma is the most widely recognized harmful eye tumor in childhood. It is portrayed by a biallelic deficiency of RB1 and is the worldview for a tumor instigated by loss of tumor silencer quality function. Because of early recognition, generally endurance of kids with retinoblastoma is high in created nations. Metastatic retinoblastoma is uncommon; however patients have a dreary visualization regardless of forceful multimodal therapy. Worldwide, less than half of patients determined to have metastatic retinoblastoma can be relieved with

current treatment conventions. Confined intraocular retinoblastoma is repairable; however numerous kids lose their vision or endure serious late impacts after forceful eye-preserving treatment. These current difficulties for pediatric oncology make the requirement for novel successful and effective restorative choices to treat these patients.

As of late, high-level articulation of the polo-like kinase (PLK1) serine/threonine kinase in retinoblastoma was related with helpless tumor separation and histopathological high-risk features. PLK1 goes about as a fundamental controller of cell cycle movement by advancing section into the M stage toward the finish of the G2 phase and prompting G2 designated spot recuperation after DNA damage. Elevated PLK1 action may animate tumor development by enacting mitotic transcriptional programs and evading the DNA-damage checkpoint. PLK1 is overexpressed in a wide assortment of malignancies, and diverse PLK1 inhibitors have been presented and examined in preclinical models and clinical preliminaries. BI6727 and GSK461364 are powerful and specific adenosine triphosphate (ATP)-competitive kinase inhibitors, which have exhibited viability against a few disease types in preclinical and clinical studies. Both inhibitors focus on the ATP-binding pocket of PLK1, official to the pivot district between the amino-terminal and the carboxy-terminal flaps of the kinase area through hydrogen connections between the dihydropteridinone inhibitor center with the spine amino and carbonyl gatherings of C133. Both inhibitors, BI6727 and GSK461364, are as monotherapy in clinical preliminaries for patients with strong tumours. BI6727 is at present being tried in blend with regular chemotherapy in a few stage II preliminaries for patients with intense myeloid leukemia and strong tumours. PLK1 inhibitors have likewise exhibited their adequacy against pediatric strong tumours, especially neuroblastomas and medulloblastomas. PLK1 downregulation has been appeared to initiate apoptosis in medulloblastoma cells and sharpen them to ionizing radiation in vitro assays. PLK1 is a potential restorative objective for retinoblastoma, and both BI6727 and GSK461364 are acceptable possibility for repressing PLK1 as a piece of tumor treatment. Here, we explored the impact of BI6727 and GSK461364 on cell suitability, cell cycle movement, apoptosis and PLK1 motioning in three cell lines got from retinoblastomas with variable hereditary foundations.

The central members controlling the cell cycle, and thusly CIN, are the numerous mitotic kinases inside the cell. These kinases drive administrative input and flagging circles that either capture the cell cycle, or drive movement through the cell cycle designated spots. Also, these kinases are regularly go about as parts of administrative criticism circles downstream of the cell cycle designated spots to guarantee obligation to cell cycle movement once designated spot necessities have been cleared. These kinases include: cyclin subordinate kinase (CDK1 or CDC2), polo-like kinase (PLK1), the Aurora kinases A, B, and C (AURKA/B/C), NIMA related kinase (NEK2), Bub1, BubR1 (or Bub1B), and TTK (or MPS1). Wee1 and Myt1 (or PKMYT1) kinases restrict the phosphatase movement of cell division cycle 25C phosphatase (CDC25C) to enact Cyclin/CDK1 buildings, yet are not viewed as mitotic kinases themselves, as they play a sign transduction job. PLK1 and CDK1 with Cyclin B are needed for going into mitosis. NEK2 action drives centrosome separation, starting the centrosome cycle. AURKA, AURKB, and PLK1 are significant for managing axle

elements and chromosome connections, with AURKA movement likewise directing focal shaft microtubule elements. AURKB likewise controls centrosome grouping for bipolar shaft development. TTK and BubR1 are basic segments of the axle get together designated spot (SAC). CDK1 and PLK1 work at the SAC to control anaphase advancing cyclosome proteasome complex (APC/C) actuation. In conclusion, PLK1 and AURKB flagging starts cytokinesis. Working in show, these kinases control a huge number of cell cycle designated spots, effector proteins, and one another and their liberation is related with various malignancies. A significant number of these atoms have been discovered to be overexpressed in tumor cells and have been advanced as possible restorative targets, including CDK1, NEK2, AURKA, Bub1, and TTK. Truth be told, our own examinations of the TCGA information across

different malignancies show their unusual articulation in different tumor types. An exact comprehension of their job in danger ought to give novel freedoms to abuse them for treatment purposes without influencing typical cells that have unblemished cell cycle designated spots. PLK1 specifically, is a central member that has been all around portrayed and with additional investigation may give significant alternatives to creating novel focused on treatments. The broad crosstalk among PLK1 and the Aurora kinases has been all around examined and PLK1 does a large number of its capacities in collaboration with the Aurora kinases, however the complexities of this relationship is looked into somewhere else, and the elements of PLK1 will be the focal point of this content. The preparing and timing of key cell cycle occasions can become dysregulated, prompting CIN through strange PLK1 exercises, and in this audit, we talk about remedial roads that may emerge from close omnipresent PLK1 liberation in disease cells.

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