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p. You can unlock the app by installing one of the unlocks on this page. It will then automatically download and install the necessary components. You can also download and install the relevant APK on any Android device, and then manually install the unlock from this page. Click the link to get access to the app. Overexpression of dual-specificity tyrosine-phosphorylation-regulated kinase in diverse hematologic malignancies. Dysregulation of cellular growth and differentiation is an important hallmark of malignancy. Indeed, dual-specificity tyrosine-phosphorylation-regulated kinase (Dyrk1a) is a ubiquitously expressed serine/threonine kinase found in normal tissues. Dyrk1a (p38, Spp38) contains a N-terminal FERM (4.1/ezrin/radixin/moesin) domain similar to the recently described kinases such as C-terminal Src kinase, focal adhesion kinase, FAK, and mitotic cyclin-dependent kinase-like kinase 1 (CDKL1). In addition, Dyrk1a has a short noncatalytic sequence at the C terminus reminiscent of catalytic domains of growth factor receptor tyrosine kinases and receptor serine/threonine kinases. Two alternatively spliced transcripts of the Dyrk1a gene, encoding two isoforms, have been described. In the present study, we cloned cDNAs encoding both isoforms of Dyrk1a. We analyzed the expression of Dyrk1a in normal and neoplastic hematopoietic cell lines and tumor samples by immunocytochemistry, Western blot analysis, and polymerase chain reaction (PCR). The expression pattern of the Dyrk1a splice variants differed between normal and malignant cells. Only the dominant, full-length isoform, Dyrk1a (p38), was expressed in normal hematopoietic cell lines and primary and malignant lymphoid cells. The expression of Dyrk1a (p38) in the B-cell lineage was elevated in Burkitt's lymphoma compared with Epstein-Barr virus-negative Burkitt's lymphoma, in Hodgkin's disease compared with normal lymphocytes, and in diffuse large cell lymphoma compared with reactive lymph nodes c6a93da74d

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