

Gene Section

Review

PRDM2 (PR domain containing 2, with ZNF domain)

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Identity

Other names: GATA 3 binding protein; HUMHOXY1; KMT8; MTB-ZF(MTE-binding protein zinc-finger); RIZ (retinoblastoma interacting zinc-finger protein); RIZ1; RIZ2
HGNC (Hugo): PRDM2
Location: 1p36.21

DNA/RNA

Description

Spans 150 Kb; 5156 bp coding sequence; 13 exons.

Transcription

Two promoters localized, respectively, in exon 1 and exon 6 guide the synthesis of two transcripts: the first one encodes for the PR domain-containing product (RIZ1) and the other one for the PR domain-lacking form (RIZ2).

Protein

Description

RIZ1 (280 kDa) and RIZ2 (260 kDa) differ only for the presence in the larger form (RIZ1) of the PR domain. In the sequence of both proteins have been identified several domains:

PR domain (130 aa), endowed with histone-methyltransferase activity (Lys 9 of histone H3) and implicated in protein-protein interactions;

E1A like-domain (100 aa), contains the LXCXE sequence able to interact with the oncosuppressor gene Rb;

Two zinc-finger clusters;
Proline-rich domain, containing a LXXLL motif mediating the interaction with estrogen receptor.

Expression

RIZ1 and RIZ2 have an ubiquitous and approxi-mately equimolar expression with higher expres-sion levels in neuroendocrine tissues.

Localisation

Nuclear.

Function

RIZ Gene products are endowed with DNA-binding as well as transcription factor-binding activities, as evidenced by the independent isolation of RIZ as a retinoblastoma-binding protein (RIZ), a DNA-binding protein (MTB-Zf), or as a GATA3 transcription factor binding protein (G3B).

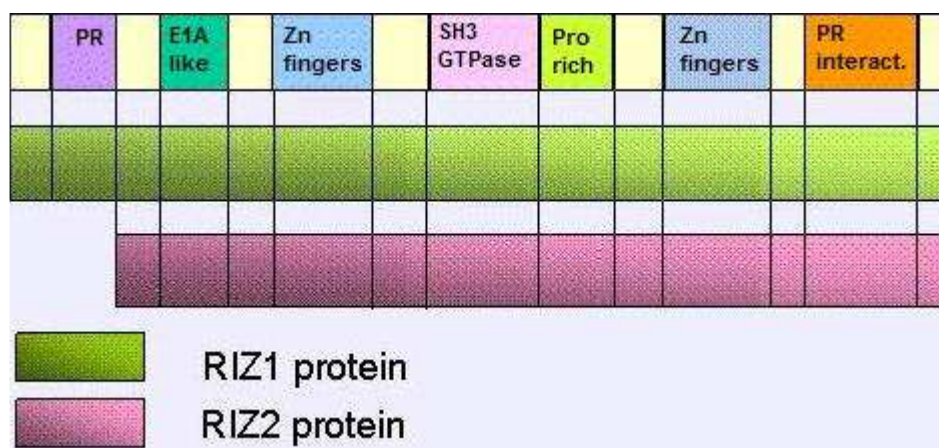
MTB-Zf (essentially identical to RIZ2) binds to the MTE DNA element GTCATATGAC of human heme-oxygenase-1 gene and can weakly activate transcription.

G3B (RIZ) interacts with the transcription factor GATA-3, regulating the expression of several genes critical for T-cell function and development.

RIZ proteins bind the estrogen receptor in a hormone-dependent manner, acting as a co-activator.

Estradiol binding to estrogen receptor complex converts RIZ activity from transcriptional repressor into co-activator.

Specific siRNA silencing of RIZ1 form increases the MCF-7 breast cancer cells growth rate. RIZ proteins act as transcriptional repressors binding to GC-rich or Sp-1-binding elements.



Mutations

Germinal

RIZ1 K.O. mice showed a high incidence of diffuse large B-cell lymphomas and a broad spectrum of unusual tumors.

Somatic

Deletion of the 1p36 region is frequent in several human cancers including neuroblastoma, breast cancer, intestinal tumors, and malignant melanoma.

Frameshift mutations in the two poly adenine tracts, A(8) and A(9), generating truncated RIZ1/2 proteins lacking the COOH-terminal PR-binding motif, were found in MSI-high (MSI-H) primary cancers occurring in the pancreas, stomach, endometrium and colorectum. The single base substitution that changed A563 in the coding sequence (immediately C-terminal to the PR domain) to a G was found specially in diffuse large B-cell lymphomas (DLBL).

Missense mutation of RIZ1 changing nucleotide G317 to A was discovered in Saos2 human osteosarcoma cells.

RIZ harbors a naturally occurring CCT insertion/deletion polymorphism in exon 8 producing a proline insertion/deletion, modulating the impact of estradiol on bone mineral density (BMD).

Implicated in

Breast cancer

Disease

The RIZ gene might have a role in human breast cancer. In fact, RIZ1 expression is decreased or lost in human breast cancer, whereas RIZ2 expression is normal.

Gastric cancer

Disease

RIZ1 might be a specific target of inactivation in human gastric cancer. Methylation of the RIZ1 promoter was detected in 37% of 30 cases and

inactivation of the second allele occurred through frameshift mutation, loss of heterozygosity (LOH) or promoter methylation. Furthermore frameshift mutations in the two coding polyadenosine tracts of RIZ were found in 19 (48%) of 40 gastric carcinomas.

Endometrial carcinoma

Disease

RIZ inactivation is highly selected during the clonal evolution of these tumors. Six (33%) of 18 endometrial carcinomas present frameshift mutation in the two coding polyadenosine tracts of RIZ.

Colorectal carcinoma

Disease

RIZ might have a role in human colorectal tumorigenesis. The region 1p36 is frequently deleted in colorectal cancer; Furthermore, 37.5% of microsatellite-unstable colorectal tumors presents frameshift mutation, consisting of 1- or 2-bp deletions of a coding (A)₈ or (A)₉ tract, of the RIZ gene.

Leukaemia

Disease

Suppression of RIZ1 expression or enhancement of RIZ2 expression may have an important role in leukaemogenesis. The expression of RIZ1 was significantly decreased in leukaemia cell lines (14 out of 17, 82%) and in patients with acute myeloblastic leukaemia (8 out of 14, 57%). In contrast, RIZ2 expression was increased in patients with acute lymphoblastic leukaemia (8 out of 11, 73%) compared with normal bone marrow cells.

Ovarian carcinoma

Disease

Reduced expression of RIZ1 may play an important role in the pathogenesis and/or development of epithelial ovarian carcinoma, and is considered to be caused in part by aberrant DNA methylation. Decreased expression of RIZ1 was significantly correlated with histological subtypes ($P < 0.0001$), high tumor grade ($P = 0.0153$), advanced clinical stage ($P = 0.0345$), and high Ki67 index ($P = 0.0117$).

Neuroblastoma

Disease

Advanced stages of neuroblastoma is genetically characterized by deletions or LOH for the short arm of chromosome 1 suggesting that in the deleted region there is a locus for putative suppressor gene.

Hepatoma

Disease

Hepatomas present deletion of the chromosome 1 short arm suggesting that in this region is localized a gene whose functional loss may be involved in hepatocellular carcinogenesis.

Prostate cancer

Disease

RIZ1 may be associated with prostate carcinogenesis. Transcriptional inactivation of RIZ1 gene by aberrant DNA methylation of promoter CpG island is detected in 20 (42.6%) of the 47 prostate cancer tissues.

Thyroid carcinoma

Disease

RIZ1 have probably an important role in thyroid tumorigenesis. RIZ1 is lost in thyroid tumor cell line and is also significantly reduced in thyroid carcinomas when compared with normal thyroid tissues and benign tumors. The loss of RIZ1 is mediated by aberrant cytosine methylation of the RIZ1 promoter.

Lung cancer

Disease

RIZ polymorphisms may be an important predictive markers for lung cancer susceptibility. In fact, the +92337G > A and +95701C > A polymorphisms are associated with reduced risk of lung adenocarcinomas.

Osteoporosis

Disease

The RIZ P704 insertion (+)/deletion (-) polymorphism modulates the impact of estradiol on bone mineral density (BMD). The RIZ P704 +/- genotype is a risk factor for low BMD in elderly subjects with low estradiol levels, whereas the RIZ P704 ++ genotype was associated with high BMD in premenopausal women.

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