

Gene Section

Review

TEK (TEK tyrosine kinase, endothelial)

Mohammad B Hossain, Nahir Cortes-Santiago, Dan Liu, Vanesa Martin, Candelaria Gomez-Manzano

Department of Neuro-Oncology, The University of Texas MD Anderson Cancer Center, Houston, TX, USA (MBH, NCS, DL, VM), Departments of Neuro-Oncology and of Genetics, The University of Texas MD Anderson Cancer Center, Houston, TX, USA (CGM)

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Identity

Other names: CD202B, CD202b, EC 2.7.10.1, hTIE2, p140 TEK, TIE-2, TIE2, VMCM, VMCM1

HGNC (Hugo): TEK

Location: 9p21.2

Note: Tie1 and Tie2 [where 'Tie' is an acronym from tyrosine kinase with Ig and EGF homology domains] are the two members of the Tie family of tyrosine kinase receptors. Tie2 has highly conserved sequence across vertebrate species, with greatest amino acid homology occurring in the kinase domain, predominantly express on the surface of endothelial cells. Three secreted natural ligands have been

characterized, - Ang1, Ang2 and interspecies orthologs Ang3 (mouse) and Ang4 (human).

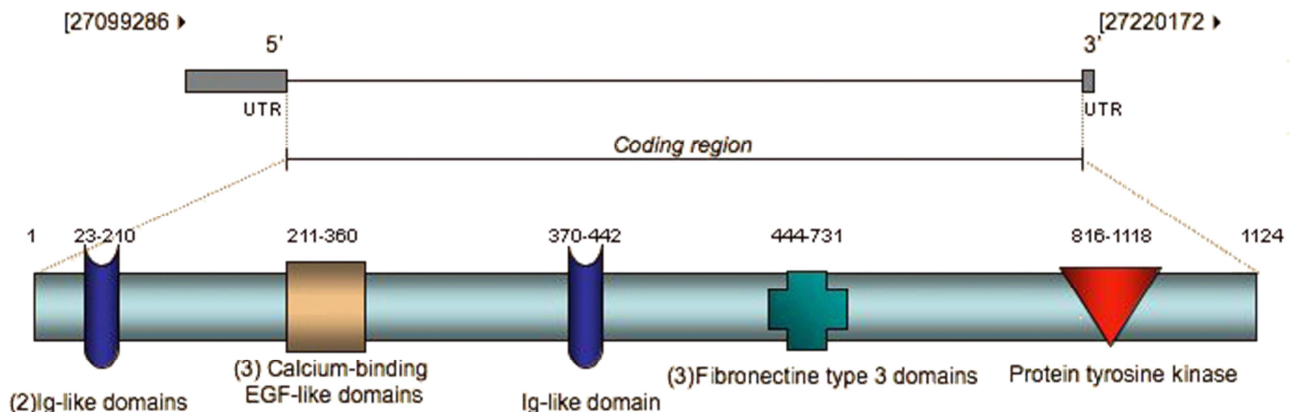
DNA/RNA

Description

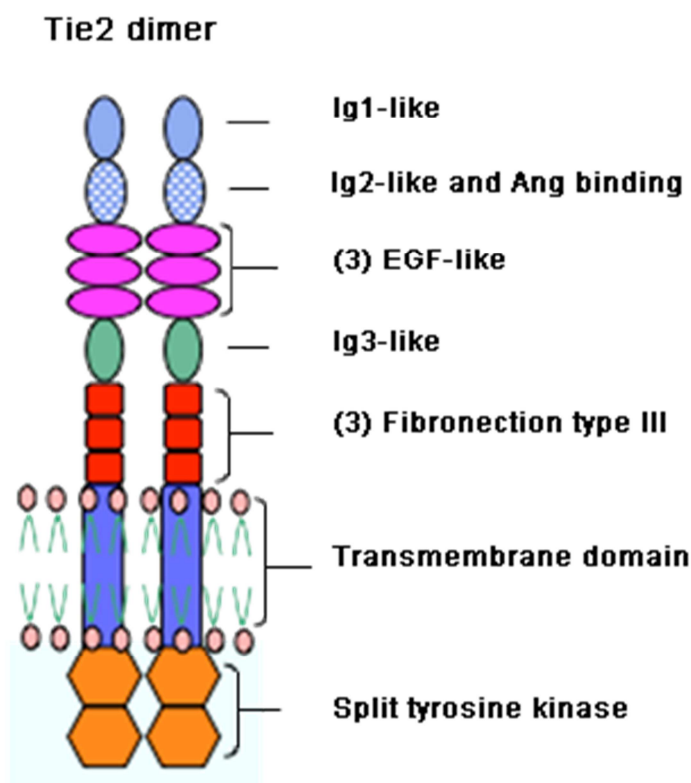
Tie2 DNA contains 121025 bp, which has 23 coding exons, in plus strand.

Transcription

mRNA contains 4817 bp transcribed in centromeric to telomeric orientation; 2 other transcripture structures by alternative splicing have been predicted. The mRNA contains a long (442 bp) 5'-UTR with 5 upstream ORFs and 1 IRES that allows the RNA to be translated under hypoxic conditions.



Schematic representation of TEK gene, coding region and protein. UTR, untranslated region; Ig, immunoglobulin; EGF, epithelial growth factor.



Protein

Description

Tie2 contains 1124 amino acids and belongs to the protein kinase superfamily, Tyr protein kinase family, Tie subfamily. Tie2 receptor contains three epidermal growth factor (EGF)-like domains flanked by three Ig-like (immunoglobulin-like) domains, followed by three fibronectin type-III domains in the extracellular domain. Tie2 possesses one intracellular split tyrosine kinase domain, which is highly conserved between the Tie members sharing a 76% sequence homology. It also contains a single transmembrane domain.

Expression

TEK/Tie2 is predominantly expressed by vascular endothelial cells: It is detectable in the critical process of new vessels formation during early development, and in the adult in response to cyclic hormonal stimulation in the ovary and uterus, as well as in healing skin wounds. High levels of expression of TEK are found in placenta, lung, spleen, and heart tissues. Hematopoietic stem cells, progenitor and mature pericytes, neural progenitor cells, and a subpopulation of monocytes (Tie2-expressing monocytes) also express Tie2/TEK. In addition, its expression might be induced in response to other pathological conditions as described below.

Localisation

Cell membrane.

Function

TEK/Tie2 is a tyrosine-kinase transmembrane receptor involved in signaling pathways upon stimuli by angiopoietins (natural ligands). It guides the proper patterning of endothelial cells during blood vessel formation and also plays role in vessel remodeling, cell survival, cell migration, and cell-to-matrix and cell-to-cell adhesion. TEK/Tie2 signaling pathway is also involved in Toll-like receptor 2 and integrin signaling.

Homology

H. sapiens: TEK;
 P. troglodytes: TEK;
 B. Taurus: TEK;
 M. musculus: Tek;
 D. rerio: tie2;
 R. norvegicus: Tek;
 G. gallus: TEK.

Mutations

Germinal

Heterozygous TEK substitutions: R849W (10 of 17 families with hereditary mucocutaneous venous malformation, VMCM), Y897S, Y897C, R915H, R918C, V1919L, A925C, K1100N. These mutations result in in vitro ligand-independent hyperphosphorylation.

Somatic

Several somatic mutations have been identified related to non-inherited vascular anomalies:

1) del-Tie2 mutant, consists in an in-frame deletion of 129bp, corresponding to a loss of exon 3 and part of exon 4 (amino acid 122-165 of extracellular Ig-like ligand-binding domain, 43 aa deletion in Ig-like domain);

2) Mutations in exon 17 of TEK/Tie2 in 49,1% (28 of 57 individuals): two somatic TEK mutations (Y897C, R915C) in vascular tumors, and seven somatic TEK mutations in vascular malformations (Y897H, Y897C, L914F, R915C, S917I, R918C, R918H).

In patients with human intramuscular haemangioma, the following Tie2 mutations have been described: G833D, Q837H.

Implicated in

Various cancers

Note

Cancer: acute myeloid leukemia and chronic myelogenous leukemia, and solid tumors, such as malignant gliomas, breast cancer, thyroid cancer, gastric cancer, bladder cancer, endometrial carcinoma. Present in both the vasculature and cancer cells of several solid tumors.

Disease

Overexpressed in the vasculature of several tumors, as breast cancer, non-small cell lung cancer, hepatocellular carcinoma, prostate cancer, Kaposi's sarcoma, and astrocytoma.

Overexpressed in the cancer cells -outside of the vascular compartment- in acute and chronic myeloid leukemia, and solid tumors, such as malignant astrocytomas, breast cancer, thyroid cancer, gastric cancer, endometrial adenocarcinoma.

Overexpressed in brain tumor stem cells and leukemic blasts.

Prognosis

Correlation of TEK/Tie2 expression and malignancy in gliomas. Tie2 activation results in increased levels of expression of ABC transporter, and eventually chemoresistance of malignant gliomas.

Tie2 regulates migration and invasion of these tumors.

Cellular Tie2 and soluble Tie2 expression might be associated to a higher risk of metastasis in patients with breast and bladder cancer, respectively.

Sporadic and inherited forms of mucocutaneous venous malformations (VMCM) and intramuscular haemangioma

Disease

Venous malformations are vascular masses composed of dilated channels lined by endothelial cells, with a

reduced coverage by pericytes, leading to functionally low resistance vessels.

See Mutation section for description of TEK/Tie2 mutations related with these diseases. Some of these mutations affect TEK/Tie2 activity and/or response to ligand.

Prognosis

Not determined.

Systemic sclerosis: microangiopathies

Disease

Systemic sclerosis (SSc) is an autoimmune disease characterized by altered angiogenesis that precede fibrosis of skin and internal organs. The abnormal angiogenesis is one of the major causes of microangiopathies.

Prognosis

Report suggesting that soluble Tie2 in serum samples from patients with systemic sclerosis is related to the development of vascular abnormalities of this disease (nailfold bleeding and pulmonary arterial hypertension).

Cytogenetics

Not determined.

Wide range of diseases with a vascular and/or inflammatory component, such as psoriasis, pulmonary hypertension, rheumatoid arthritis

Disease

Psoriatic lesions are characterized by elongation and dilatation of papillary dermis. Tie2 transgenic mice showed epidermal hyperproliferation, inflammatory cell accumulation, and altered dermal angiogenesis, mimicking the phenotype present in human psoriasis.

Pulmonary hypertension is characterized by high pulmonary arterial pressure. Increased levels of activated TEK/Tie2 and Angiotensin II have been reported.

Rheumatoid arthritis is characterized by chronic inflammatory changes in synovial tissue and pathological angiogenesis. Tie2 has been shown to be upregulated in synovial tissue of patients with rheumatoid arthritis and to mediate pathological angiogenesis and invasion within affected synovial tissue.

Prognosis

Not determined.

Cytogenetics

Not determined.

Cerebrovascular disease: stroke

Disease

In stroke, arterial occlusion of one or more arteries in the brain leads to focal or global ischemia and tissue damage and eventual death. Expression of activated

Tie2 and Ang1 lead to enhanced neovascularization, vessel stabilization and improved recovery.

Prognosis

Reports suggest that activation of Tie2-mediated pathway is essential in initiation of survival responses in neural progenitor cells against cerebral ischemia and hypoxia.

Cytogenetics

Not determined.

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