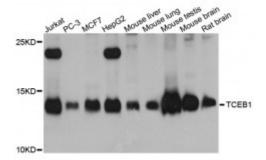


Anti-TCEB1 Antibody





Description This gene encodes the protein elongin C, which is a subunit of the

transcription factor B (SIII) complex. The SIII complex is composed of elongins A/A2, B and C. It activates elongation by RNA polymerase II by suppressing transient pausing of the polymerase at many sites within transcription units. Elongin A functions as the transcriptionally active component of the SIII complex, whereas elongins B and C are regulatory subunits. Elongin A2 is specifically expressed in the testis, and capable of forming a stable complex with elongins B and C. The von Hippel-Lindau tumor suppressor protein binds to elongins B and C, and thereby inhibits transcription elongation. Multiple alternatively spliced transcript variants encoding two distinct isoforms have been identified.

Model STJ114389

Host Rabbit

Reactivity Human, Mouse, Rat

Applications WB

Immunogen Recombinant fusion protein containing a sequence corresponding to amino

acids 1-112 of human TCEB1 (NP_001191791.1).

Gene ID 6921

Gene Symbol <u>ELOC</u>

Dilution range WB 1:500 - 1:2000

Tissue Specificity Overexpressed in prostate cancer cell line PC-3 and breast cancer cell line SK-

BR-3

Purification Affinity purification

Note For Research Use Only (RUO).

Protein Name Elongin-C EloC Elongin 15 kDa subunit RNA polymerase II transcription

factor SIII subunit C SIII p15 Transcription elongation factor B polypeptide 1

Molecular Weight 12.473 kDa

Clonality Polyclonal

Conjugation Unconjugated

Isotype IgG

Formulation PBS with 0.02% sodium azide, 50% glycerol, pH7.3.

Storage Instruction Store at -20C. Avoid freeze / thaw cycles.

Database Links HGNC:11617OMIM:600788Reactome:R-HSA-112382

Alternative Names Elongin-C EloC Elongin 15 kDa subunit RNA polymerase II transcription

factor SIII subunit C SIII p15 Transcription elongation factor B polypeptide 1

Function SIII, also known as elongin, is a general transcription elongation factor that

increases the RNA polymerase II transcription elongation past template-

encoded arresting sites, Subunit A is transcriptionally active and its

transcription activity is strongly enhanced by binding to the dimeric complex of the SIII regulatory subunits B and C (elongin BC complex), By binding to BC-box motifs it seems to link target recruitment subunits, like VHL and members of the SOCS box family, to Cullin/RBX1 modules that activate E2

ubiquitination enzymes,

Cellular Localization Nucleus

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