

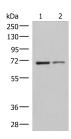
Tel:240-252-7368(USA) Fax: 240-252-7376(USA) techsupport@elabscience.com Website: www.elabscience.com

EWSR1 Polyclonal Antibody

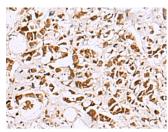
Catalog No.E-AB-52908ReactivityH,MStorageStore at -20°C. Avoid freeze / thaw cycles.HostRabbitApplicationsWB,IHC,ELISAIsotypeIgG

Note: Centrifuge before opening to ensure complete recovery of vial contents.

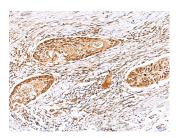
Images



Western blot analysis of K562 and HepG2 cell lysates using EWSR1 Polyclonal Antibody at dilution of 1:1000



Immunohistochemistry of paraffinembedded Human breast cancer tissue using EWSR1 Polyclonal Antibody at dilution of 1:60(×200)



Immunohistochemistry of paraffinembedded Human esophagus cancer tissue using EWSR1 Polyclonal Antibody at dilution of 1:60(×200)

Immunogen Information

Immunogen Fusion protein of human EWSR1

Gene Accession BC000527 **Swissprot** Q01844

Synonyms EWS,EWS oncogene,EWS RNA binding protein

1,EWS,EWSR 1,Ewsr1,EWSR1 protein

Product Information

Calculated MW 68 kDa

Observed MW Refer to figures

Buffer PBS with 0.05% NaN3 and 40% Glycerol,pH7.4

Purify Antigen affinity purification

Dilution WB 1:1000-1:5000, IHC 1:50-1:300, ELISA

1:5000-1:10000

Background

This gene encodes a multifunctional protein that is involved in various cellular processes, including gene expression, cell signaling, and RNA processing and transport. The protein includes an N-terminal transcriptional activation domain and a C-terminal RNA-binding domain. Chromosomal translocations between this gene and various genes encoding transcription factors result in the production of chimeric proteins that are involved in tumorigenesis. These chimeric proteins usually consist of the N-terminal transcriptional activation domain of this protein fused to the C-terminal DNA-binding domain of the transcription factor protein. Mutations in this gene, specifically a t(11;22)(q24;q12) translocation, are known to cause Ewing sarcoma as well as neuroectodermal and various other tumors. Alternative splicing of this gene results in multiple transcript variants. Related pseudogenes have been identified on chromosomes 1 and 14.

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