EWSR1 (Human) Recombinant Protein (Q01)

Catalog # H00002130-Q01 Size 25 ug, 10 ug

Applications



Specification	
Product Description	Human EWSR1 partial ORF (NP_005234, 358 a.a 453 a.a.) recombinant protein with GST-tag at N-terminal.
Sequence	SDNSAIYVQGLNDSVTLDDLADFFKQCGVVKMNKRTGQPMIHIYLDKETGKPKGDATVSYEDPPT AKAAVEWFDGKDFQGSKLKVSLARKKPPMNS
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	36.3
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCI, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.
Storage Instruction	Store at -80°C. Aliquot to avoid repeated freezing and thawing.
Note	Best use within three months from the date of receipt of this protein.

Applications

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- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — EWSR1	
Entrez GenelD	2130
GeneBank Accession#	<u>NM_005243</u>
Protein Accession#	<u>NP_005234</u>
Gene Name	EWSR1
Gene Alias	EWS
Gene Description	Ewing sarcoma breakpoint region 1
Omim ID	<u>133450</u>
Gene Ontology	Hyperlink
Gene Summary	This gene encodes a multifunctional protein that is involved in various cellular processes, includin g gene expression, cell signaling, and RNA processing and transport. The protein includes an N-t erminal transcriptional activation domain and a C-terminal RNA-binding domain. Chromosomal tr anslocations between this gene and various genes encoding transcription factors result in the pro duction of chimeric proteins that are involved in tumorigenesis. These chimeric proteins usually co nsist of the N-terminal transcriptional activation domain of this protein fused to the C-terminal DN A-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 vario us other tumors. Alternative splicing of this gene results in multiple transcript variants. Related pse udogenes have been identified on chromosomes 1 and 14. [provided by RefSeq
Other Designations	Ewings sarcoma EWS-Fli1 (type 1) oncogene bK984G1.4 (Ewing sarcoma breakpoint region 1 p