

# DHX8 Antibody (monoclonal) (M07)

Mouse monoclonal antibody raised against a partial recombinant DHX8. Catalog # AT1761a

### **Product Information**

**Application** WB, E **Primary Accession** Q14562 **Other Accession** NM 004941 Reactivity Human Host mouse Clonality monoclonal Isotype IgG1 Kappa **Clone Names** 1D6 **Calculated MW** 139315

### **Additional Information**

**Gene ID** 1659

Other Names ATP-dependent RNA helicase DHX8, DEAH box protein 8, RNA helicase HRH1,

DHX8, DDX8

Target/Specificity DHX8 (NP\_004932, 301 a.a. ~ 400 a.a) partial recombinant protein with GST

tag. MW of the GST tag alone is 26 KDa.

**Dilution** WB~~1:500~1000 E~~N/A

**Format** Clear, colorless solution in phosphate buffered saline, pH 7.2.

**Storage** Store at -20°C or lower. Aliquot to avoid repeated freezing and thawing.

**Precautions** DHX8 Antibody (monoclonal) (M07) is for research use only and not for use in

diagnostic or therapeutic procedures.

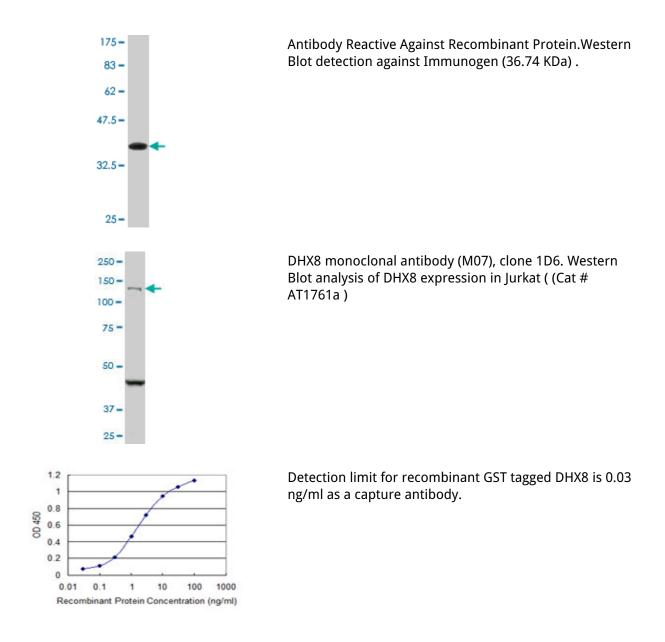
## **Background**

DEAD box proteins, characterized by the conserved motif Asp-Glu-Ala-Asp (DEAD), are putative RNA helicases. They are implicated in a number of cellular processes involving alteration of RNA secondary structure such as translation initiation, nuclear and mitochondrial splicing, and ribosome and spliceosome assembly. Based on their distribution patterns, some members of this family are believed to be involved in embryogenesis, spermatogenesis, and cellular growth and division. This gene encodes a DEAD box protein, which is highly homologous to yeast Prp22. This protein facilitates nuclear export of spliced mRNA by releasing the RNA from the spliceosome.

### References

Crystallization and preliminary X-ray diffraction analysis of the C-terminal domain of the human spliceosomal DExD/H-box protein hPrp22. Kudlinzki D, et al. Acta Crystallogr Sect F Struct Biol Cryst Commun, 2009 Sep 1. PMID 19724143. Defining the human deubiquitinating enzyme interaction landscape. Sowa ME, et al. Cell, 2009 Jul 23. PMID 19615732. Novel function of beta-arrestin2 in the nucleus of mature spermatozoa. Neuhaus EM, et al. J Cell Sci, 2006 Aug 1. PMID 16820410. Quantitative phosphoproteome analysis using a dendrimer conjugation chemistry and tandem mass spectrometry. Tao WA, et al. Nat Methods, 2005 Aug. PMID 16094384. Nucleolar proteome dynamics. Andersen JS, et al. Nature, 2005 Jan 6. PMID 15635413.

### **Images**



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