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## **Datasheet**

## TAF11 monoclonal antibody (M01), clone 3D3

Catalog Number: H00006882-M01

Regulatory Status: For research use only (RUO)

**Product Description:** Mouse monoclonal antibody

raised against a partial recombinant TAF11.

Clone Name: 3D3

 $\label{eq:local_local_local_local} \begin{tabular}{ll} \textbf{Immunogen:} TAF11 & (NP\_005634, 158 a.a. $\sim 210 a.a) \\ \textbf{partial recombinant protein with GST tag.} & MW of the \\ \end{tabular}$ 

GST tag alone is 26 KDa.

Sequence:

SKVFVGEVVEEALDVCEKWGEMPPLQPKHMREAVRR LKSKGQIPNSKHKKIIF

Host: Mouse

Reactivity: Human

Applications: ELISA, IF, IHC-P, S-ELISA, WB-Ce,

WB-Re

(See our web site product page for detailed applications

information)

Protocols: See our web site at

http://www.abnova.com/support/protocols.asp or product

page for detailed protocols

Isotype: IgG1 Kappa

Storage Buffer: In 1x PBS, pH 7.4

Storage Instruction: Store at -20°C or lower. Aliquot to

avoid repeated freezing and thawing.

Entrez GenelD: 6882

Gene Symbol: TAF11

Gene Alias: MGC:15243, PRO2134, TAF2I, TAFII28

**Gene Summary:** Initiation of transcription by RNA polymerase II requires the activities of more than 70 polypeptides. The protein that coordinates these activities is transcription factor IID (TFIID), which binds

to the core promoter to position the polymerase properly, serves as the scaffold for assembly of the remainder of the transcription complex, and acts as a channel for regulatory signals. TFIID is composed of the TATA-binding protein (TBP) and a group of evolutionarily conserved proteins known as TBP-associated factors or TAFs. TAFs may participate in basal transcription, serve as coactivators, function in promoter recognition or modify general transcription factors (GTFs) to facilitate complex assembly and transcription initiation. This gene encodes a small subunit of TFIID that is present in all TFIID complexes and interacts with TBP. This subunit also interacts with another small subunit, TAF13, to form a heterodimer with a structure similar to the histone core structure. [provided by RefSeq]