

## YTHDF2 Antibody - C-terminal region

Rabbit Polyclonal Antibody Catalog # Al15546

### **Specification**

# YTHDF2 Antibody - C-terminal region - Product Information

Application WB
Primary Accession Other Accession NP 057342

Reactivity Human, Mouse,

Rat, Rabbit, Horse, Bovine,

Dog

Predicted Human, Mouse,

Rat, Rabbit, Horse, Bovine,

Host Rabbit
Clonality Polyclonal
Calculated MW 62kDa KDa

YTHDF2 Antibody - C-terminal region - Additional Information

### **Gene ID 51441**

# Alias Symbol HGRG8, NY-REN-2 Other Names

YTH domain-containing family protein 2, CLL-associated antigen KW-14,

High-glucose-regulated protein 8, Renal carcinoma antigen NY-REN-2, YTHDF2, HGRG8

#### **Format**

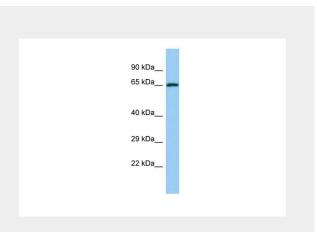
Liquid. Purified antibody supplied in 1x PBS buffer with 0.09% (w/v) sodium azide and 2% sucrose.

#### Reconstitution & Storage

Add 50 ul of distilled water. Final anti-YTHDF2 antibody concentration is 1 mg/ml in PBS buffer with 2% sucrose. For longer periods of storage, store at 20°C. Avoid repeat freeze-thaw cycles.

### **Precautions**

YTHDF2 Antibody - C-terminal region is for research use only and not for use in



Host: Rabbit

Target Name: YTHDF2

Sample Tissue: Jurkat Whole cell lysate

S

Antibody Dilution: 1.0µg/mlYTHDF2 is

supported by BioGPS gene expression data to

be expressed in Jurkat

# YTHDF2 Antibody - C-terminal region - References

Scanlan M.J., et al.Int. J. Cancer 83:456-464(1999).

Roberts T.P., et al. Submitted (OCT-1999) to the EMBL/GenBank/DDBI databases.

LINDL/Geribarik/DDbj databases.

Krackhardt A.M., et al. Submitted (OCT-2001) to the EMBL/GenBank/DDBI databases.

Ota T., et al. Nat. Genet. 36:40-45(2004).

Gregory S.G., et al. Nature 441:315-321(2006).



diagnostic or therapeutic procedures.

YTHDF2 Antibody - C-terminal region - Protein Information

Name YTHDF2

{ECO:0000303|PubMed:24284625, ECO:0000312|HGNC:HGNC:31675}

#### **Function**

Specifically recognizes and binds N6-methyladenosine (m6A)- containing RNAs, and regulates their stability (PubMed:<a href="http://www.uniprot.org/citations/24284625"

target="\_blank">24284625</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/26046440"

target=" blank">26046440</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/26318451"

target=" blank">26318451</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/32492408"

target="\_blank">32492408</a>). M6A is a modification present at internal sites of mRNAs and some non-coding RNAs and plays a role in mRNA stability and processing (PubMed:<a href="http://www.u">http://www.u</a>

niprot.org/citations/22575960"

target="\_blank">22575960</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/24284625"

target=" blank">24284625</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/32492408"

target="\_blank">32492408</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/25412658"

target=" blank">25412658</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/25412661"

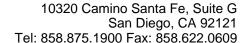
target="\_blank">25412661</a>). Acts as a regulator of mRNA stability by promoting degradation of m6A-containing mRNAs via interaction with the CCR4-NOT and ribonuclease P/MRP complexes, depending on the context (PubMed:<a href="http://www.uniprot.org/citations/24284625" target="\_blank">24284625</a>,

PubMed: <a href="http://www.uniprot.org/ci tations/26046440"

target="\_blank">26046440</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/27558897"

target=" blank">27558897</a>,

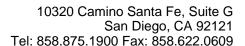




PubMed:<a href="http://www.uniprot.org/ci tations/30930054" target="\_blank">30930054</a>, PubMed:<a href="http://www.uniprot.org/ci tations/32492408" target=" blank">32492408</a>). The YTHDF paralogs (YTHDF1, YTHDF2 and YTHDF3) share m6A-containing mRNAs targets and act redundantly to mediate mRNA degradation and cellular differentiation (PubMed:<a href="http://ww w.uniprot.org/citations/28106072" target=" blank">28106072</a>, PubMed:<a href="http://www.uniprot.org/ci tations/32492408" target=" blank">32492408</a>). M6A-containing mRNAs containing a binding site for RIDA/HRSP12 (5'-GGUUC-3') are preferentially degraded by endoribonucleolytic cleavage: cooperative binding of RIDA/HRSP12 and YTHDF2 to transcripts leads to recruitment of the ribonuclease P/MRP complex (PubMed:<a hr ef="http://www.uniprot.org/citations/30930 054" target=" blank">30930054</a>). Other m6A-containing mRNAs undergo deadenylation via direct interaction between YTHDF2 and CNOT1, leading to recruitment of the CCR4-NOT and subsequent deadenylation of m6Acontaining mRNAs (PubMed: <a href="http:/ /www.uniprot.org/citations/27558897" target=" blank">27558897</a>). Required maternally to regulate oocyte maturation: probably acts by binding to m6A-containing mRNAs, thereby regulating maternal transcript dosage during oocyte maturation, which is essential for the competence of oocytes to sustain early zygotic development (By similarity). Also required during spermatogenesis: regulates spermagonial adhesion by promoting degradation of m6A-containing transcripts coding for matrix metallopeptidases (By similarity). Also involved in hematopoietic stem cells specification by binding to m6A-containing mRNAs, leading to promote their degradation (PubMed:<a href="http:// www.uniprot.org/citations/30065315" target=" blank">30065315</a>). Also acts as a regulator of neural development by promoting m6A-dependent degradation of neural development-related mRNA targets (By similarity). Inhibits neural specification of induced pluripotent stem cells by binding to methylated neural-specific mRNAs and promoting their degradation, thereby



restraining neural differentiation (PubMed:<a href="http://www.uniprot.org/c itations/32169943" target="\_blank">32169943</a>). Regulates circadian regulation of hepatic lipid metabolism: acts by promoting m6A-dependent degradation of PPARA transcripts (PubMed:<a href="http://www.u niprot.org/citations/30428350" target="\_blank">30428350</a>). Regulates the innate immune response to infection by inhibiting the type I interferon response: acts by binding to m6A-containing IFNB transcripts and promoting their degradation (PubMed:<a hr ef="http://www.uniprot.org/citations/30559" 377" target="\_blank">30559377</a>). May also act as a promoter of cap-independent mRNA translation following heat shock stress: upon stress, relocalizes to the nucleus and specifically binds mRNAs with some m6A methylation mark at their 5'-UTR, protecting demethylation of mRNAs by FTO, thereby promoting cap-independent mRNA translation (PubMed:<a href="http://www.u niprot.org/citations/26458103" target=" blank">26458103</a>). Regulates mitotic entry by promoting the phase-specific m6A-dependent degradation of WEE1 transcripts (PubMed:<a href="http" ://www.uniprot.org/citations/32267835" target=" blank">32267835</a>). Promotes formation of phase-separated membraneless compartments, such as P-bodies or stress granules, by undergoing liquid-liquid phase separation upon binding to mRNAs containing multiple m6A-modified residues: polymethylated mRNAs act as a multivalent scaffold for the binding of YTHDF proteins, juxtaposing their disordered regions and thereby leading to phase separation (PubMed:<a href="http:// www.uniprot.org/citations/31388144" target=" blank">31388144</a>, PubMed: <a href="http://www.uniprot.org/ci tations/31292544" target=" blank">31292544</a>, PubMed:<a href="http://www.uniprot.org/ci tations/32451507" target=" blank">32451507</a>, PubMed: <a href="http://www.uniprot.org/ci"> tations/31642031" target=" blank">31642031</a>). The resulting mRNA-YTHDF complexes then partition into different endogenous phase-separated membraneless





compartments, such as P-bodies, stress granules or neuronal RNA granules (PubMed:<a href="http://www.uniprot.org/c itations/31292544" target="\_blank">31292544</a>). May also recognize and bind RNAs modified by C5-methylcytosine (m5C) and act as a regulator of rRNA processing (PubMed:<a h ref="http://www.uniprot.org/citations/31815440" target="\_blank">31815440</a>).

#### **Cellular Location**

Cytoplasm, cytosol. Cytoplasm, P-body. Cytoplasm, Stress granule. Nucleus. Note=Localizes to the cytosol and relocates to the nucleus following heat shock stress (PubMed:26458103) Can partition into different structures: into P-bodies in unstressed cells, and into stress granules during stress (PubMed:31292544)

### **Tissue Location**

Highly expressed in induced pluripotent stem cells (iPSCs) and down-regulated during neural differentiation

# YTHDF2 Antibody - C-terminal region - Protocols

Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- <u>Immunohistochemistry</u>
- Immunofluorescence
- Immunoprecipitation
- Flow Cytomety
- Cell Culture