

### QKI antibody - N-terminal region

Rabbit Polyclonal Antibody Catalog # Al15031

#### **Specification**

### **QKI antibody - N-terminal region - Product Information**

Application WB
Primary Accession Q96PU8

Other Accession <u>NM 206853</u>, <u>NP 996735</u>

Reactivity Human, Mouse, Rat, Rabbit, Pig, Horse,

**Bovine, Guinea Pig, Dog** 

Predicted Human, Mouse, Rat, Pig, Chicken, Horse,

**Bovine, Guinea Pig, Dog** 

Host Rabbit
Clonality Polyclonal
Calculated MW 35kDa KDa

# **QKI antibody - N-terminal region - Additional Information**

**Gene ID 9444** 

Alias Symbol DKFZp586I0923, Hqk, QK, QK1, QK3, hqkl Other Names

Protein quaking, Hqk, Hqkl, QKI, HKQ

#### 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-QKI 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**

QKI antibody - N-terminal region is for research use only and not for use in diagnostic or therapeutic procedures.

### QKI antibody - N-terminal region - Protein Information

Name QKI {ECO:0000303|PubMed:16342280, ECO:0000312|HGNC:HGNC:21100}

#### **Function**

RNA reader protein, which recognizes and binds specific RNAs, thereby regulating RNA metabolic processes, such as pre-mRNA splicing, circular RNA (circRNA) formation, mRNA export, mRNA stability and/or translation (PubMed:<a href="http://www.uniprot.org/citations/22398723" target="\_blank">22398723</a>, PubMed:<a href="http://www.uniprot.org/citations/23630077" target="\_blank">23630077</a>, PubMed:<a href="http://www.uniprot.org/citations/25768908" target="\_blank">25768908</a>, PubMed:<a href="http://www.uniprot.org/citations/27029405" target="\_blank">27029405</a>, PubMed:<a href="http://www.uniprot.org/citations/31331967"



target="\_blank">31331967</a>, PubMed:<a href="http://www.uniprot.org/citations/37379838" target="blank">37379838</a>). Involved in various cellular processes, such as mRNA storage into stress granules, apoptosis, lipid deposition, interferon response, glial cell fate and development (PubMed: <a href="http://www.uniprot.org/citations/25768908" target=" blank">25768908</a>, PubMed:<a href="http://www.uniprot.org/citations/31829086" target=" blank">31829086</a>, PubMed:<a href="http://www.uniprot.org/citations/34428287" target=" blank">34428287</a>, PubMed:<a href="http://www.uniprot.org/citations/37379838" target="blank">37379838</a>). Binds to the 5'-NACUAAY-N(1,20)-UAAY-3' RNA core sequence (PubMed:<a href="http://www.uniprot.org/citations/23630077" target=" blank">23630077</a>). Acts as a mRNA modification reader that specifically recognizes and binds mRNA transcripts modified by internal N(7)-methylguanine (m7G) (PubMed:<a href="http://www.uniprot.org/citations/37379838" target=" blank">37379838</a>). Promotes the formation of circular RNAs (circRNAs) during the epithelial to mesenchymal transition and in cardiomyocytes: acts by binding to sites flanking circRNA-forming exons (PubMed: <a href="http://www.uniprot.org/citations/25768908" target=" blank">25768908</a>). CircRNAs are produced by back-splicing circularization of pre-mRNAs (PubMed:<a href="http://www.uniprot.org/citations/25768908" target=" blank">25768908</a>). Plays a central role in myelinization via 3 distinct mechanisms (PubMed:<a href="http://www.uniprot.org/citations/16641098" target=" blank">16641098</a>). First, acts by protecting and promoting stability of target mRNAs such as MBP, SIRT2 and CDKN1B, which promotes oligodendrocyte differentiation (By similarity). Second, participates in mRNA transport by regulating the nuclear export of MBP mRNA (By similarity). Finally, indirectly regulates mRNA splicing of MAG pre- mRNA during oligodendrocyte differentiation by acting as a negative regulator of MAG exon 12 alternative splicing; acts by binding to HNRNPA1 mRNA splicing factor, preventing its translation (By similarity). Involved in microglia differentiation and remyelination by regulating microexon alternative splicing of the Rho GTPase pathway (By similarity). Involved in macrophage differentiation: promotes monocyte differentiation by regulating pre-mRNA splicing in naive peripheral blood monocytes (PubMed:<a href="http://www.uniprot.org/citations/27029405" target=" blank">27029405</a>). Acts as an important regulator of muscle development: required for the contractile function of cardiomyocytes by regulating alternative splicing of cardiomyocyte transcripts (By similarity). Acts as a negative regulator of thermogenesis by decreasing stability, nuclear export and translation of mRNAs encoding PPARGC1A and UCP1 (By similarity). Also required for visceral endoderm function and blood vessel development (By similarity). May also play a role in smooth muscle development (PubMed: <a href="http://www.uniprot.org/citations/31331967" target=" blank">31331967</a>). In addition to its RNA-binding activity, also acts as a nuclear transcription coactivator for SREBF2/SREBP2 (By similarity).

### **Cellular Location**

Nucleus. Cytoplasm [Isoform QKI6]: Cytoplasm, cytosol. Nucleus Note=Localizes predominantly in the cytoplasm and at lower levels in nucleus.

#### **Tissue Location**

Expressed in the frontal cortex of brain. Down-regulated in the brain of schizophrenic patients

#### **QKI antibody - N-terminal region - Protocols**

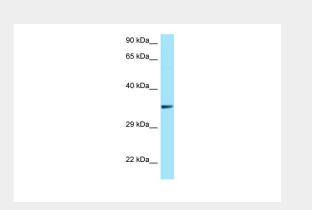
Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- <u>Immunofluorescence</u>
- <u>Immunoprecipitation</u>



- Flow Cytomety
- Cell Culture

# QKI antibody - N-terminal region - Images



WB Suggested Anti-QKI Antibody Titration: 1.0 μg/ml

Positive Control: Hela Whole CellThere is BioGPS gene expression data showing that QKI is expressed in HeLa

# **QKI antibody - N-terminal region - References**

Li Z.Z., et al. Jpn. J. Cancer Res. 93:167-177(2002).

Xia J.-H., et al. Submitted (APR-1999) to the EMBL/GenBank/DDBJ databases.

Li H., et al. Submitted (DEC-2005) to the EMBL/GenBank/DDBJ databases.

Mungall A.J., et al. Nature 425:805-811(2003).

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