

# **DGCR8 / Pasha Antibody (internal region)**

Peptide-affinity purified goat antibody Catalog # AF2905a

## **Specification**

## DGCR8 / Pasha Antibody (internal region) - Product Information

Application

Primary Accession <u>O8WYQ5</u>

Other Accession NP 073557.3, 54487, 94223 (mouse), 287954

<u>(rat)</u>

Predicted Human, Mouse, Rat, Cow

Host Goat
Clonality Polyclonal
Concentration 0.5 mg/ml
Isotype IgG

Calculated MW 86045

# DGCR8 / Pasha Antibody (internal region) - Additional Information

#### **Gene ID 54487**

## **Other Names**

Microprocessor complex subunit DGCR8, DiGeorge syndrome critical region 8, DGCR8, C22orf12, DGCRK6

#### **Format**

0.5 mg/ml in Tris saline, 0.02% sodium azide, pH7.3 with 0.5% bovine serum albumin

#### Storage

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

## **Precautions**

DGCR8 / Pasha Antibody (internal region) is for research use only and not for use in diagnostic or therapeutic procedures.

## DGCR8 / Pasha Antibody (internal region) - Protein Information

### Name DGCR8

Synonyms C22orf12, DGCRK6

#### **Function**

Component of the microprocessor complex that acts as a RNA- and heme-binding protein that is involved in the initial step of microRNA (miRNA) biogenesis. Component of the microprocessor complex that is required to process primary miRNA transcripts (pri-miRNAs) to release precursor miRNA (pre-miRNA) in the nucleus. Within the microprocessor complex, DGCR8 function as a



molecular anchor necessary for the recognition of pri-miRNA at dsRNA-ssRNA junction and directs DROSHA to cleave 11 bp away form the junction to release hairpin-shaped pre-miRNAs that are subsequently cut by the cytoplasmic DICER to generate mature miRNAs (PubMed: <a href="http://www.uniprot.org/citations/26027739" target="\_blank">26027739</a>, PubMed:<a href="http://www.uniprot.org/citations/26748718" target="\_blank">26748718</a>). The hemebound DGCR8 dimer binds pri-miRNAs as a cooperative trimer (of dimers) and is active in triggering pri-miRNA cleavage, whereas the heme-free DGCR8 monomer binds pri-miRNAs as a dimer and is much less active. Both double-stranded and single-stranded regions of a pri-miRNA are required for its binding (PubMed: <a href="http://www.uniprot.org/citations/15531877" target=" blank">15531877</a>, PubMed:<a href="http://www.uniprot.org/citations/15574589" target="blank">15574589</a>, PubMed:<a href="http://www.uniprot.org/citations/15589161" target="blank">15589161</a>, PubMed:<a href="http://www.uniprot.org/citations/16751099" target="blank">16751099</a>, PubMed:<a href="http://www.uniprot.org/citations/16906129" target="blank">16906129</a>, PubMed:<a href="http://www.uniprot.org/citations/16963499" target="blank">16963499</a>, PubMed:<a href="http://www.uniprot.org/citations/17159994" target="blank">17159994</a>). Specifically recognizes and binds N6-methyladenosine (m6A)-containing pri-miRNAs, a modification required for pri-miRNAs processing (PubMed: <a href="http://www.uniprot.org/citations/25799998" target="\_blank">25799998</a>). Involved in

#### **Cellular Location**

Nucleus. Nucleus, nucleolus. Note=Colocalizes with nucleolin and DROSHA in the nucleolus. Mostly detected in the nucleolus as electron- dense granular patches around the fibrillar center (FC) and granular component (GC). Also detected in the nucleoplasm as small foci adjacent to splicing speckles near the chromatin structure. Localized with DROSHA in GW bodies (GWBs), also known as P-bodies (PubMed:17159994)

**Tissue Location** 

Ubiquitously expressed.

# DGCR8 / Pasha Antibody (internal region) - Protocols

the silencing of embryonic stem cell self-renewal (By similarity).

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

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

## DGCR8 / Pasha Antibody (internal region) - Images

# DGCR8 / Pasha Antibody (internal region) - References

Altered brain microRNA biogenesis contributes to phenotypic deficits in a 22q11-deletion mouse model. Stark KL, Xu B, Bagchi A, Lai WS, Liu H, Hsu R, Wan X, Pavlidis P, Mills AA, Karayiorgou M, Gogos JA. Nat. Genet. 2008 Jun 40 (6): 751-60. PMID: 18469815