bs-12450R

[Primary Antibody]

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AGO2 Rabbit pAb

DATASHEET -

Host: Rabbit Isotype: IgG

Clonality: Polyclonal

GenelD: 27161 SWISS: Q9UKV8

Target: AGO2

Immunogen: KLH conjugated synthetic peptide derived from human AGO2:

431-530/859.

Purification: affinity purified by Protein A

Concentration: 1mg/ml

Storage: 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50%

Glycerol.

Shipped at 4°C. Store at -20°C for one year. Avoid repeated

freeze/thaw cycles.

Background: Eukaryotic translation initiation factor 2C (eIF2C) proteins (argonaute family) influence RNA interference (RNAi) as components of the RNA-inducible silencing complex (RISC) or microRNA (miRNA)-containing ribonucleoprotein particle (miRNP). Small RNAs, including small interfering RNAs (siRNAs) and miRNAs, can silence target genes through mechanisms that utilize RISC or miRNP particles. eIF2C1 (argonaute 1, AGO1, eIF2C, GERP95, Q99) and Dicer1 play a coordinated role in siRNA-mediated gene silencing. eIF2C2 (Slicer, argonaute 2, AGO2, Q10) is a RISC component that can concentrate in cytoplasmic processing bodies (P-bodies) and catalyze mRNA cleavage. Mammalian P-bodies contain mRNAs and have an association with miRNA-induced translational silencing and siRNA-induced mRNA degradation. Additional eIF2C proteins include eIF2C3 (argonaute 3, AGO3), elF2C4 (argonaute 4, AGO4) and melF2c5 (mouse argonaute 5).

Applications: WB (1:500-2000)

IHC-P (1:100-500) **IHC-F** (1:100-500) **IF** (1:100-500)

Reactivity: Mouse (predicted: Human,

Rat, Pig, Sheep, Cow, Dog,

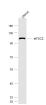
Horse)

Predicted 97 kDa

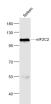
MW.:

Subcellular Location: Cytoplasm ,Nucleus

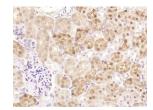
VALIDATION IMAGES



Sample: Uterus (Mouse) Lysate at 40 ug Primary: Anti-eIF2C2 (bs-12450R) at 1/1000 dilution Secondary: IRDve800CW Goat Anti-Rabbit IgG at 1/20000 dilution Predicted band size: 97 kD Observed band size: 97 kD



Sample: Spleen (Mouse) Lysate at 40 ug Primary: Anti-eIF2C2 (bs-12450R) at 1/1000 dilution Secondary: IRDye800CW Goat Anti-Rabbit IgG at 1/20000 dilution Predicted band size: 97 kD Observed band size: 97 kD



Paraformaldehyde-fixed, paraffin embedded (mouse kidney); Antigen retrieval by boiling in sodium citrate buffer (pH6.0) for 15min; Block endogenous peroxidase by 3% hydrogen peroxide for 20 minutes; Blocking buffer (normal goat serum) at 37°C for 30min; Antibody incubation with (AGO2) Polyclonal Antibody, Unconjugated (bs-12450R) at 1:200 overnight at 4°C, followed by operating according to SP Kit(Rabbit) (sp-0023) instructions and DAB staining.

— SELECTED CITATIONS -

- [IF=11.161] Zhipeng Jiang, et al. EIF4A3-induced circ_0084615 contributes to the progression of colorectal cancer via miR-599/ONECUT2 pathway. J Exp Clin Canc Res. 2021 Dec;40(1):1-15 IP; Human. 34253241
- [IF=6.208] Xiaoyv Yue. et al. circITGB1 Regulates Adipocyte Proliferation and Differentiation via the miR-23a/ARRB1

- Pathway. INT J MOL SCI. 2023 Jan;24(3):1976 Other; Sheep. 36768295
- [IF=4.5] Duo Cheng. et al. Downregulation of circ-RAPGEF5 inhibits colorectal cancer progression by reducing the expression of polypeptide N-acetylgalactosaminyltransferase 3 (GALNT3). ENVIRON TOXICOL. 2024 May;: Other ;Human. 38775215
- [IF=4.657] Feng, Lei. et al. Circ_0088194 Regulates Proliferation, Migration, Apoptosis, and Inflammation by miR-30a-3p/ADAM10 Axis in Rheumatoid Arthritis Fibroblastic Synovial Cells. INFLAMMATION. 2022 Aug;:1-14 Other ;Human. 35918573
- [IF=3.223] Wenshu Yang. et al. Circular RNA hsa_circ_0008003 promotes the progression of non-small-cell lung cancer by sponging miR-548I and regulating KPNA4 expression. THORAC CANCER. 2022 Dec;: Other; Human. 36573325