bsm-52419R

[Primary Antibody]

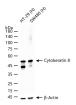
Cytokeratin 8 Recombinant Rabbit mAb



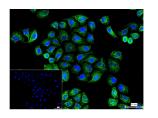
www.bioss.com.cn sales@bioss.com.cn techsupport@bioss.com.cn 400-901-9800

| – DATASHEET ––––– | | 400-901-9800 |
|---|--------------------|---|
| Host: Rabbit | Isotype: IgG | Applications: WB (1:1000-5000) |
| Clonality: Recombina | Int CloneNo.: 8C12 | IHC-P (1:200-800) IHC-F (1:200-800) |
| GenelD: 3856 | SWISS: P05787 | IF (1:200-800) |
| Target: Cytokeratin | ۱8 | ICC/IF (1:100-500) |
| Immunogen: A synthesized peptide derived from human Cytokeratin 8: 450-483. | | 8. Reactivity: Human, Mouse, Rat |
| Purification: affinity puri | ified by Protein A | |
| Concentration: 1mg/ml | | |
| Storage: 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol. Store at -20°C for one year. Avoid repeated freeze/thaw cycles. The lyophilized antibody is stable at room temperature for at least one month and for greater than a year when kept at -20°C. When reconstituted in sterile pH 7.4 0.01M PBS or diluent of antibody the antibody is stable for at least two weeks at 2-4°C. | | e Subcellular Location: Cytoplasm ,Nucleus |
| Background: This gene is a member of the type II keratin family clustered on the long arm of chromosome 12. Type I and type II keratins heteropolymerize to form intermediate-sized filaments in the cytoplasm of epithelial cells. The product of this gene typically dimerizes with keratin 18 to form an intermediate filament in simple single-layered epithelial cells. This protein plays a role in maintaining cellular structural integrity and also functions in signal transduction and cellular differentiation. Mutations in this gene cause cryptogenic cirrhosis. Alternatively spliced transcript variants have been found for this gene. [provided by RefSeq, Jan 2012]. | | |

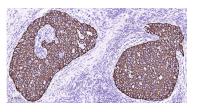
— VALIDATION IMAGES -



25 ug total protein per lane of various lysates (see on figure) probed with Cytokeratin 8 monoclonal antibody, unconjugated (bsm-52419R) at 1:1000 dilution and 4°C overnight incubation. Followed by conjugated secondary antibody incubation at r.t. for 60 min.

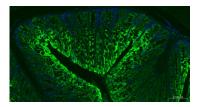


4% Paraformaldehyde-fixed Hela (H) cell; Triton X-100 at r.t. for 20 min; Antibody incubation with (Cytokeratin 8) monoclonal Antibody, unconjugated (bsm-52419R) 1:100, 90 min at

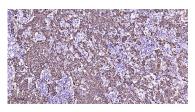


Paraformaldehyde-fixed, paraffin embedded Human Breast Cancer; Antigen retrieval by boiling in sodium citrate buffer (pH6.0) for 15 min; Antibody incubation with Cytokeratin 8 Monoclonal Antibody,

 $\label{eq:unconjugated} Unconjugated(bsm-52419R) at 1:200 overnight at 4°C, followed by conjugation to the bs-0295G-HRP and DAB (C-0010) staining.$

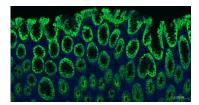


Paraformaldehyde-fixed, paraffin embedded Mouse colon; Antigen retrieval by boiling in sodium citrate buffer (pH6.0) for 15 min; Antibody incubation with Cytokeratin 8



Paraformaldehyde-fixed, paraffin embedded Human Thyroid Cancer; Antigen retrieval by boiling in sodium citrate buffer (pH6.0) for 15 min; Antibody incubation with Cytokeratin 8 Monoclonal Antibody,

Unconjugated(bsm-52419R) at 1:200 overnight at 4°C, followed by conjugation to the bs-0295G-HRP and DAB (C-0010) staining.



Paraformaldehyde-fixed, paraffin embedded Human stomach; Antigen retrieval by boiling in sodium citrate buffer (pH6.0) for 15 min; Antibody incubation with Cytokeratin 8

37°C; followed by conjugated Goat Anti-Rabbit IgG antibody (green, bs-40295G-FITC) at 37°C for 90 min, DAPI (blue, C02-04002) was used to stain the cell nuclei. PBS instead of the primary antibody was used as the blank control. Monoclonal Antibody, Unconjugated (bsm-52419R) at 1:200 overnight at 4°C. Followed by conjugated Goat Anti-Rabbit IgG antibody (green, bs-0295G-BF488), DAPI (blue, C02-04002) was used to stain the cell nuclei. Monoclonal Antibody, Unconjugated (bsm-52419R) at 1:200 overnight at 4°C. Followed by conjugated Goat Anti-Rabbit IgG antibody (green, bs-0295G-BF488), DAPI (blue, C02-04002) was used to stain the cell nuclei.

- SELECTED CITATIONS -

• [IF=6.8] Ye Jiazhou. et al. Single cell-spatial transcriptomics and bulk multi-omics analysis of heterogeneity and ecosystems in hepatocellular carcinoma. NPJ PRECIS ONCOL. 2024 Nov;8(1):1-18 IHC ;Human. 39548284