

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 9, 2026

normal goat IgG

RRID:AB_737167

Type: Antibody

Proper Citation

Santa Cruz Biotechnology Cat# sc-2028, RRID:AB_737167

Antibody Information

URL: http://antibodyregistry.org/AB_737167

Proper Citation: Santa Cruz Biotechnology Cat# sc-2028, RRID:AB_737167

Target Antigen: normal goat IgG

Host Organism: goat

Clonality: unknown

Comments: validation status unknown check with seller; recommendations: IgG

Antibody Name: normal goat IgG

Description: This unknown targets normal goat IgG

Antibody ID: AB_737167

Vendor: Santa Cruz Biotechnology

Catalog Number: sc-2028

Record Creation Time: 20250820T051022+0000

Record Last Update: 20250820T205902+0000

Ratings and Alerts

- ENCODE PROJECT External validation for lot: B0311 is available under ENCODE ID: ENCAB462PUJ - ENCODE
<https://www.encodeproject.org/antibodies/ENCAB462PUJ>

No alerts have been found for normal goat IgG.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 22 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Akhtar F, et al. (2026) The RNA-binding protein HuR modulates the expression of the disease-linked CCL2 rs1024611G-rs13900T haplotype. eLife, 13.

Kawamura S, et al. (2025) Modeling antithymocyte globulin-induced microvasculopathy using human iPSC-derived vascularized liver organoids. Cell reports. Medicine, 6(11), 102433.

Nagao M, et al. (2025) Chromatin remodeler Chd7 regulates reactive astrogliosis after ischemic stroke. Cell reports, 44(12), 116590.

Kim J, et al. (2024) An enhancer RNA recruits KMT2A to regulate transcription of Myb. Cell reports, 43(7), 114378.

Sekiya M, et al. (2023) Loss of CtBP2 may be a mechanistic link between metabolic derangements and progressive impairment of pancreatic β cell function. Cell reports, 42(8), 112914.

Abe K, et al. (2022) Distinct patterns of RNA polymerase II and transcriptional elongation characterize mammalian genome activation. Cell reports, 41(13), 111865.

Li X, et al. (2022) β -Defensin 19/119 mediates sperm chemotaxis and is associated with idiopathic infertility. Cell reports. Medicine, 3(12), 100825.

Cheung VC, et al. (2021) Pluripotent stem cell-derived endometrial stromal fibroblasts in a cyclic, hormone-responsive, coculture model of human decidua. Cell reports, 35(7), 109138.

Andreeva L, et al. (2021) NLRP3 cages revealed by full-length mouse NLRP3 structure control pathway activation. Cell, 184(26), 6299.

Li Z, et al. (2021) Acetyl-CoA Synthetase 2: A Critical Linkage in Obesity-Induced Tumorigenesis in Myeloma. Cell metabolism, 33(1), 78.

Fujii K, et al. (2021) Controlling tissue patterning by translational regulation of signaling transcripts through the core translation factor eIF3c. Developmental cell, 56(21), 2928.

Bi Y, et al. (2020) Identification of ALPPL2 as a Naive Pluripotent State-Specific Surface Protein Essential for Human Naive Pluripotency Regulation. Cell reports, 30(11), 3917.

McDaniel MM, et al. (2020) Suppression of Inflammasome Activation by IRF8 and IRF4 in cDCs Is Critical for T Cell Priming. Cell reports, 31(5), 107604.

Jiang L, et al. (2020) Bach1-induced suppression of angiogenesis is dependent on the BTB domain. *EBioMedicine*, 51, 102617.

Crosby P, et al. (2019) Insulin/IGF-1 Drives PERIOD Synthesis to Entrain Circadian Rhythms with Feeding Time. *Cell*, 177(4), 896.

Boonying W, et al. (2019) Pink1 regulates FKBP5 interaction with AKT/PHLPP and protects neurons from neurotoxin stress induced by MPP. *Journal of neurochemistry*, 150(3), 312.

Davis MR, et al. (2019) Nuclear eIF4E Stimulates 3'-End Cleavage of Target RNAs. *Cell reports*, 27(5), 1397.

Minegishi T, et al. (2018) Shootin1b Mediates a Mechanical Clutch to Produce Force for Neuronal Migration. *Cell reports*, 25(3), 624.

Zhu Z, et al. (2017) PHB Associates with the HIRA Complex to Control an Epigenetic-Metabolic Circuit in Human ESCs. *Cell stem cell*, 20(2), 274.

Dave K, et al. (2017) Mice deficient of Myc super-enhancer region reveal differential control mechanism between normal and pathological growth. *eLife*, 6.