

Resource Summary Report

Generated by [RRID](#) on Jul 29, 2026

IRDye 680LT Goat anti-Rabbit IgG

RRID:AB_10706309

Type: Antibody

Proper Citation

LICORbio Cat# 926-68021, RRID:AB_10706309

Antibody Information

URL: http://antibodyregistry.org/AB_10706309

Proper Citation: LICORbio Cat# 926-68021, RRID:AB_10706309

Target Antigen: IgG

Host Organism: goat

Clonality: unknown

Comments: Applications: Western blotting

Info: Reacts with the heavy and light chains of rabbit IgG, and with the light chains of rabbit IgM and IgA.

Antibody Name: IRDye 680LT Goat anti-Rabbit IgG

Description: This unknown targets IgG

Target Organism: rabbit

Antibody ID: AB_10706309

Vendor: LICORbio

Catalog Number: 926-68021

Record Creation Time: 20250820T050431+0000

Record Last Update: 20250820T204306+0000

Ratings and Alerts

No rating or validation information has been found for IRDye 680LT Goat anti-Rabbit IgG.

No alerts have been found for IRDye 680LT Goat anti-Rabbit IgG.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 54 mentions in open access literature.

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

Arimura Y, et al. (2025) MagIC-Cryo-EM, structural determination on magnetic beads for scarce macromolecules in heterogeneous samples. *eLife*, 13.

Atwa A, et al. (2025) BioID2-Based Tau Interactome Reveals Novel and Known Protein Interactions Associated with Multiple Cellular Pathways. *Journal of proteome research*.

Flaherty JN, et al. (2025) The catalytic efficiency of METTL16 affects cellular processes by governing the intracellular S-adenosylmethionine setpoint. *Cell reports*, 44(7), 115966.

Sun Y, et al. (2025) TBK1 Targeting Is Identified as a Therapeutic Strategy to Enhance CAR T-Cell Efficacy Using Patient-Derived Organotypic Tumor Spheroids. *Cancer immunology research*, 13(2), 210.

Hofman DA, et al. (2024) Translation of non-canonical open reading frames as a cancer cell survival mechanism in childhood medulloblastoma. *Molecular cell*, 84(2), 261.

Ugur B, et al. (2024) VPS13B is localized at the interface between Golgi cisternae and is a functional partner of FAM177A1. *The Journal of cell biology*, 223(12).

Bayazitov IT, et al. (2024) Sound-evoked adenosine release in cooperation with neuromodulatory circuits permits auditory cortical plasticity and perceptual learning. *Cell reports*, 43(2), 113758.

DaRosa PA, et al. (2024) UFM1 E3 ligase promotes recycling of 60S ribosomal subunits from the ER. *Nature*, 627(8003), 445.

Achom M, et al. (2024) A genetic basis for sex differences in Xp11 translocation renal cell carcinoma. *Cell*, 187(20), 5735.

Serebrenik YV, et al. (2024) Pooled endogenous protein tagging and recruitment for systematic profiling of protein function. *Cell genomics*, 4(10), 100651.

Sabbarini IM, et al. (2023) Zinc-finger protein Zpr1 is a bespoke chaperone essential for eEF1A biogenesis. *Molecular cell*, 83(2), 252.

Dark N, et al. (2023) Generation of left ventricle-like cardiomyocytes with improved structural, functional, and metabolic maturity from human pluripotent stem cells. *Cell reports methods*, 3(4), 100456.

McQuown AJ, et al. (2023) A Zpr1 co-chaperone mediates folding of eukaryotic translation elongation factor 1A via a GTPase cycle. *Molecular cell*, 83(17), 3108.

Wu L, et al. (2023) CD28-CAR-T cell activation through FYN kinase signaling rather than LCK enhances therapeutic performance. *Cell reports. Medicine*, 4(2), 100917.

Valenti M, et al. (2023) Human gasdermin D and MLKL disrupt mitochondria, endocytic traffic and TORC1 signalling in budding yeast. *Open biology*, 13(5), 220366.

Hofman DA, et al. (2023) Translation of non-canonical open reading frames as a cancer cell survival mechanism in childhood medulloblastoma. *bioRxiv : the preprint server for biology*.

Schmit KJ, et al. (2023) Fiber deprivation and microbiome-borne curli shift gut bacterial populations and accelerate disease in a mouse model of Parkinson's disease. *Cell reports*, 42(9), 113071.

Tang S, et al. (2022) A genome-scale CRISPR screen reveals PRMT1 as a critical regulator of androgen receptor signaling in prostate cancer. *Cell reports*, 38(8), 110417.

Kim M, et al. (2022) YAP1 and PRDM14 converge to promote cell survival and tumorigenesis. *Developmental cell*, 57(2), 212.

Rigter PMF, et al. (2022) Adult Camk2a gene reinstatement restores the learning and plasticity deficits of Camk2a knockout mice. *iScience*, 25(11), 105303.