Resource Summary Report

Generated by dkNET on Sep 1, 2026

Anti-Glutathione-S-Transferase (GST) antibody produced in rabbit

RRID:AB_259965
Type: Antibody

Proper Citation

Sigma-Aldrich Cat# G7781, RRID:AB_259965

Antibody Information

URL: http://antibodyregistry.org/AB_259965

Proper Citation: Sigma-Aldrich Cat# G7781, RRID:AB_259965

Target Antigen: Glutathione-S-Transferase (GST) antibody produced in rabbit

Host Organism: rabbit

Clonality: polyclonal

Comments: Vendor recommendations: ELISA; Other; Western Blot; indirect ELISA: suitable immunoblotting: 1:2,000 using purified recombinant GST or lysates of induced Escherichia coli expressing GST fusion proteins, dot blot: suitable

Antibody Name: Anti-Glutathione-S-Transferase (GST) antibody produced in rabbit

Description: This polyclonal targets Glutathione-S-Transferase (GST) antibody produced in rabbit

Antibody ID: AB_259965

Vendor: Sigma-Aldrich

Catalog Number: G7781

Record Creation Time: 20250820T005135+0000

Record Last Update: 20250820T092340+0000

Ratings and Alerts

No rating or validation information has been found for Anti-Glutathione-S-Transferase (GST) antibody produced in rabbit.

No alerts have been found for Anti-Glutathione-S-Transferase (GST) antibody produced in rabbit.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 30 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Jagannatha P, et al. (2026) Large-scale tethered screen of RNA-binding proteins reveals novel regulators of poly(A) site selection. Molecular cell, 86(14), 2811.

Vind AC, et al. (2026) Mechanism of RACK1-dependent ZAK? activation at stalled and collided ribosomes. Molecular cell, 86(12), 2341.

Zhao XT, et al. (2025) The Myo2 adaptor Ldm1 and its receptor Ldo16 mediate actin-dependent lipid droplet motility. Cell reports, 44(11), 116475.

Lambourne L, et al. (2025) Widespread variation in molecular interactions and regulatory properties among transcription factor isoforms. Molecular cell, 85(7), 1445.

Qin X, et al. (2025) Light-quality-directed plant growth strategy controlled by SnRK2s. Developmental cell.

King J, et al. (2025) Missense variants in human forkhead transcription factors reveal determinants of forkhead DNA bispecificity. Cell reports, 44(11), 116427.

Wang J, et al. (2024) ARF alters PAF1 complex integrity to selectively repress oncogenic transcription programs upon p53 loss. Molecular cell, 84(23), 4538.

Szczesna M, et al. (2024) Bacterial esterases reverse lipopolysaccharide ubiquitylation to block host immunity. Cell host & microbe, 32(6), 913.

Toda T, et al. (2024) Mrc1 regulates parental histone segregation and heterochromatin inheritance. Molecular cell, 84(17), 3223.

Grochowska KM, et al. (2023) Jacob-induced transcriptional inactivation of CREB promotes A?-induced synapse loss in Alzheimer's disease. The EMBO journal, 42(4), e112453.

Kapoor S, et al. (2023) PP2A-B55SUR-6 promotes nuclear envelope breakdown in C. elegans embryos. Cell reports, 42(12), 113495.

Gao Z, et al. (2023) A pair of readers of bivalent chromatin mediate formation of Polycomb-based "memory of cold" in plants. Molecular cell, 83(7), 1109.

Curtis AJ, et al. (2023) Molecular basis of interactions between CaMKII and ?-actinin-2 that underlie dendritic spine enlargement. eLife, 12.

Wang L, et al. (2023) TONSOKU is required for the maintenance of repressive chromatin modifications in Arabidopsis. Cell reports, 42(7), 112738.

Xie W, et al. (2022) CYLD deubiquitinates plakoglobin to promote Cx43 membrane targeting and gap junction assembly in the heart. Cell reports, 41(13), 111864.

Zang Y, et al. (2021) GhUBX controlling helical growth results in production of stronger cotton fiber. iScience, 24(8), 102930.

Studniarek C, et al. (2021) The 7SK/P-TEFb snRNP controls ultraviolet radiation-induced transcriptional reprogramming. Cell reports, 35(2), 108965.

Bacon CW, et al. (2020) KAP1 Is a Chromatin Reader that Couples Steps of RNA Polymerase II Transcription to Sustain Oncogenic Programs. Molecular cell, 78(6), 1133.

Catania S, et al. (2020) Evolutionary Persistence of DNA Methylation for Millions of Years after Ancient Loss of a De Novo Methyltransferase. Cell, 180(2), 263.

Wu W, et al. (2020) The Autophagy-Initiating Kinase ULK1 Controls RIPK1-Mediated Cell Death. Cell reports, 31(3), 107547.