

# Resource Summary Report

Generated by [NIF](#) on Aug 27, 2026

## Stat2 (C-20)

RRID:AB\_632437

Type: Antibody

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### Proper Citation

Santa Cruz Biotechnology Cat# sc-476, RRID:AB\_632437

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_632437](http://antibodyregistry.org/AB_632437)

**Proper Citation:** Santa Cruz Biotechnology Cat# sc-476, RRID:AB\_632437

**Target Antigen:** Stat2 (C-20)

**Host Organism:** rabbit

**Clonality:** polyclonal

**Comments:** Discontinued: 2016; validation status unknown check with seller; recommendations: Immunohistochemistry; Western Blot; Immunoprecipitation; Immunofluorescence; Immunocytochemistry; ELISA; WB, IP, IF, IHC(P), ELISA

**Antibody Name:** Stat2 (C-20)

**Description:** This polyclonal targets Stat2 (C-20)

**Target Organism:** human

**Antibody ID:** AB\_632437

**Vendor:** Santa Cruz Biotechnology

**Catalog Number:** sc-476

**Record Creation Time:** 20250820T010048+0000

**Record Last Update:** 20250820T094747+0000

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### Ratings and Alerts

- ENCODE PROJECT External validation for lot: B1908 is available under ENCODE ID: ENCAB000ALG - ENCODE

<https://www.encodeproject.org/antibodies/ENCAB000ALG>

**Warning:** Discontinued: 2016

Discontinued: 2016; validation status unknown check with seller; recommendations: Immunohistochemistry; Western Blot; Immunoprecipitation; Immunofluorescence; Immunocytochemistry; ELISA; WB, IP, IF, IHC(P), ELISA

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## Data and Source Information

**Source:** [Antibody Registry](#)

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Kehrer T, et al. (2023) Impact of SARS-CoV-2 ORF6 and its variant polymorphisms on host responses and viral pathogenesis. Cell host & microbe, 31(10), 1668.

Le-Trilling VTK, et al. (2020) The Human Cytomegalovirus pUL145 Isoforms Act as Viral DDB1-Cullin-Associated Factors to Instruct Host Protein Degradation to Impede Innate Immunity. Cell reports, 30(7), 2248.

Shah PS, et al. (2018) Comparative Flavivirus-Host Protein Interaction Mapping Reveals Mechanisms of Dengue and Zika Virus Pathogenesis. Cell, 175(7), 1931.