

# Resource Summary Report

Generated by [kravitz2](#) on Aug 11, 2026

## Rabbit anti-UBAP2L/NICE4 Antibody, Affinity Purified

RRID:AB\_477953

Type: Antibody

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### Proper Citation

Bethyl Cat# A300-533A, RRID:AB\_477953

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_477953](http://antibodyregistry.org/AB_477953)

**Proper Citation:** Bethyl Cat# A300-533A, RRID:AB\_477953

**Target Antigen:** UBAP2L/NICE4

**Host Organism:** rabbit

**Clonality:** polyclonal

**Comments:** Applications: WB, IP  
Original Manufacturer

**Antibody Name:** Rabbit anti-UBAP2L/NICE4 Antibody, Affinity Purified

**Description:** This polyclonal targets UBAP2L/NICE4

**Target Organism:** human

**Antibody ID:** AB\_477953

**Vendor:** Bethyl

**Catalog Number:** A300-533A

**Alternative Catalog Numbers:** A300-533A-M, A300-533A-T

**Record Creation Time:** 20250820T061824+0000

**Record Last Update:** 20250820T234522+0000

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### Ratings and Alerts

- ENCODE PROJECT External validation for lot: 1 is available under ENCODE ID: ENCAB201KCT - ENCODE  
<https://www.encodeproject.org/antibodies/ENCAB201KCT>

**Warning:** Discontinued at Thermo Fisher Scientific  
Applications: WB, IP  
Original Manufacturer

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Antal CE, et al. (2023) A super-enhancer-regulated RNA-binding protein cascade drives pancreatic cancer. Nature communications, 14(1), 5195.

Wheeler EC, et al. (2020) Pooled CRISPR screens with imaging on microraft arrays reveals stress granule-regulatory factors. Nature methods, 17(6), 636.

Zhu G, et al. (2020) TRIM11 Prevents and Reverses Protein Aggregation and Rescues a Mouse Model of Parkinson's Disease. Cell reports, 33(9), 108418.

Markmiller S, et al. (2019) Active Protein Neddylation or Ubiquitylation Is Dispensable for Stress Granule Dynamics. Cell reports, 27(5), 1356.

Youn JY, et al. (2018) High-Density Proximity Mapping Reveals the Subcellular Organization of mRNA-Associated Granules and Bodies. Molecular cell, 69(3), 517.