## **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on May 6, 2025

# hnRNP C1/C2 (4F4)

RRID:AB\_627731 Type: Antibody

#### **Proper Citation**

(Santa Cruz Biotechnology Cat# sc-32308, RRID:AB\_627731)

### **Antibody Information**

URL: http://antibodyregistry.org/AB\_627731

**Proper Citation:** (Santa Cruz Biotechnology Cat# sc-32308, RRID:AB\_627731)

Target Antigen: Human HNRNPC

**Host Organism:** mouse

Clonality: monoclonal

**Comments:** validation status unknown check with seller; recommendations: Immunofluorescence; Immunoprecipitation; Western Blot; Western Blotting,

Immunoprecipitation, Immunofluorescence

Antibody Name: hnRNP C1/C2 (4F4)

**Description:** This monoclonal targets Human HNRNPC

Target Organism: human

Clone ID: 4F4

Antibody ID: AB\_627731

Vendor: Santa Cruz Biotechnology

Catalog Number: sc-32308

**Record Creation Time:** 20231110T043810+0000

Record Last Update: 20241115T110847+0000

#### **Ratings and Alerts**

No rating or validation information has been found for hnRNP C1/C2 (4F4).

No alerts have been found for hnRNP C1/C2 (4F4).

#### Data and Source Information

Source: Antibody Registry

#### **Usage and Citation Metrics**

We found 8 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch.org.

Zhang X, et al. (2024) Multivalent GU-rich oligonucleotides sequester TDP-43 in the nucleus by inducing high molecular weight RNP complexes. iScience, 27(6), 110109.

Cheng Y, et al. (2024) A non-canonical role for a small nucleolar RNA in ribosome biogenesis and senescence. Cell, 187(17), 4770.

Ghanam AR, et al. (2023) Alternative transcribed 3' isoform of long non-coding RNA Malat1 inhibits mouse retinal oxidative stress. iScience, 26(1), 105740.

Miao W, et al. (2023) Glucose dissociates DDX21 dimers to regulate mRNA splicing and tissue differentiation. Cell, 186(1), 80.

Pan Y, et al. (2023) Extracellular Vesicle-Mediated Transfer of LncRNA IGFL2-AS1 Confers Sunitinib Resistance in Renal Cell Carcinoma. Cancer research, 83(1), 103.

Krismer K, et al. (2020) Transite: A Computational Motif-Based Analysis Platform That Identifies RNA-Binding Proteins Modulating Changes in Gene Expression. Cell reports, 32(8), 108064.

Lee SH, et al. (2019) Gain of Additional BIRC3 Protein Functions through 3'-UTR-Mediated Protein Complex Formation. Molecular cell, 74(4), 701.

Nozawa RS, et al. (2017) SAF-A Regulates Interphase Chromosome Structure through Oligomerization with Chromatin-Associated RNAs. Cell, 169(7), 1214.