

Resource Summary Report

Generated by [RRID](#) on Aug 1, 2026

Anti-Bax, NT

RRID:AB_310143

Type: Antibody

Proper Citation

Millipore Cat# 06-499, RRID:AB_310143

Antibody Information

URL: http://antibodyregistry.org/AB_310143

Proper Citation: Millipore Cat# 06-499, RRID:AB_310143

Target Antigen: Bax NT

Host Organism: rabbit

Clonality: polyclonal

Comments: seller recommendations: IgG; IgG IC, IH, IH(P), IP, WB; Immunocytochemistry; Western Blot; Immunohistochemistry; Immunoprecipitation

Antibody Name: Anti-Bax, NT

Description: This polyclonal targets Bax NT

Target Organism: h, m

Antibody ID: AB_310143

Vendor: Millipore

Catalog Number: 06-499

Record Creation Time: 20250820T010813+0000

Record Last Update: 20250820T100738+0000

Ratings and Alerts

No rating or validation information has been found for Anti-Bax, NT.

No alerts have been found for Anti-Bax, NT.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Dengler MA, et al. (2021) BAX mitochondrial integration is regulated allosterically by its ? 1-?2 loop. Cell death and differentiation, 28(12), 3270.

Doerflinger M, et al. (2020) Flexible Usage and Interconnectivity of Diverse Cell Death Pathways Protect against Intracellular Infection. Immunity, 53(3), 533.

Iyer S, et al. (2020) Robust autoactivation for apoptosis by BAK but not BAX highlights BAK as an important therapeutic target. Cell death & disease, 11(4), 268.

Lin KH, et al. (2019) Systematic Dissection of the Metabolic-Apoptotic Interface in AML Reveals Heme Biosynthesis to Be a Regulator of Drug Sensitivity. Cell metabolism, 29(5), 1217.

Dengler MA, et al. (2019) BAX Activation: Mutations Near Its Proposed Non-canonical BH3 Binding Site Reveal Allosteric Changes Controlling Mitochondrial Association. Cell reports, 27(2), 359.

Iyer S, et al. (2019) Probing BAK and BAX Activation and Pore Assembly with Cytochrome c Release, Limited Proteolysis, and Oxidant-Induced Linkage. Methods in molecular biology (Clifton, N.J.), 1877, 201.

Robin AY, et al. (2018) Ensemble Properties of Bax Determine Its Function. Structure (London, England : 1993), 26(10), 1346.