

Resource Summary Report

Generated by NIF on Jul 29, 2026

PE anti-human CD127 (IL-7R?)

RRID:AB_2564136

Type: Antibody

Proper Citation

BioLegend Cat# 351340, RRID:AB_2564136

Antibody Information

URL: http://antibodyregistry.org/AB_2564136

Proper Citation: BioLegend Cat# 351340, RRID:AB_2564136

Target Antigen: CD127

Host Organism: mouse

Clonality: monoclonal

Comments: Applications: FC

Antibody Name: PE anti-human CD127 (IL-7R?)

Description: This monoclonal targets CD127

Target Organism: human

Clone ID: Clone A019D5

Antibody ID: AB_2564136

Vendor: BioLegend

Catalog Number: 351340

Alternative Catalog Numbers: 351304, 351303

Record Creation Time: 20250820T000249+0000

Record Last Update: 20250820T071234+0000

Ratings and Alerts

No rating or validation information has been found for PE anti-human CD127 (IL-7R?).

No alerts have been found for PE anti-human CD127 (IL-7R?).

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Dyikanov D, et al. (2024) Comprehensive peripheral blood immunoprofiling reveals five immunotypes with immunotherapy response characteristics in patients with cancer. Cancer cell, 42(5), 759.

Barra JM, et al. (2024) Combinatorial genetic engineering strategy for immune protection of stem cell-derived beta cells by chimeric antigen receptor regulatory T cells. Cell reports, 43(11), 114994.

Li Z, et al. (2024) Therapeutic application of human type 2 innate lymphoid cells via induction of granzyme B-mediated tumor cell death. Cell, 187(3), 624.

Forsthuber A, et al. (2024) Cancer-associated fibroblast subtypes modulate the tumor-immune microenvironment and are associated with skin cancer malignancy. Nature communications, 15(1), 9678.

Li Z, et al. (2024) Isolation, expansion, and adoptive transfer of human ILC2s for the treatment of mice bearing liquid and solid tumors. STAR protocols, 5(3), 103096.

Zaitsev A, et al. (2022) Precise reconstruction of the TME using bulk RNA-seq and a machine learning algorithm trained on artificial transcriptomes. Cancer cell, 40(8), 879.